GenCore version 5.1.6

OM nucleic - nucleic search, using sw model

Run on: January 21, 2004, 14:51:06 ; Search time 1041 Seconds
(without alignments)
11139.742 Million cell updates/sec

Title: US-09-981-353-172
Perfect score: 3290
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Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 2324096 seqs, 1762381658 residues

Total number of hits satisfying chosen parameters: 4648192

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_NA:*
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18: /cgn2_6/ptodata/2/pubpna/US60_PUBCOMB.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	3290	100.0	3290	10	US-09-981-353-172	Sequence 172, App
2	3290	100.0	3290	11	US-09-919-039-164	Sequence 164, App
3	3270.8	99.4	3328	13	US-10-240-965-111	Sequence 111, App
4	3175.2	96.5	3220	10	US-09-954-531-137	Sequence 137, App
5	3175.2	96.5	3220	10	US-09-954-531-356	Sequence 356, App
6	3175.2	96.5	3220	12	US-10-314-790-2	Sequence 2, Appli
7	3175.2	96.5	3220	13	US-10-021-660-3	Sequence 3, Appli
8	3175.2	96.5	3220	13	US-09-873-367C-694	Sequence 694, App
9	2911.2	88.5	2916	13	US-10-024-298A-124	Sequence 124, App
10	2911.2	88.5	2916	13	US-10-042-211A-124	Sequence 124, App
11	2245.8	68.3	3447	12	US-10-062-674-2102	Sequence 2102, Ap
12	617.2	18.8	3175	12	US-10-314-790-4	Sequence 4, Appli
13	617.2	18.8	3175	13	US-09-873-367C-143	Sequence 143, App
14	615	18.7	2287	12	US-10-388-934-200	Sequence 200, App
15	495.2	15.1	662	10	US-09-879-536-623	Sequence 623, App
16	459.8	14.0	463	10	US-09-867-701-2803	Sequence 2803, Ap
17	451.4	13.7	613	10	US-09-879-536-521	Sequence 521, App
c 18	448	13.6	495	10	US-09-920-300A-1304	Sequence 1304, Ap
c 19	448	13.6	495	13	US-10-099-926-1304	Sequence 1304, Ap
c 20	448	13.6	495	14	US-10-033-528-1304	Sequence 1304, Ap
21	430	13.1	441	10	US-09-867-701-2808	Sequence 2808, Ap
c 22	415	12.6	2747	13	US-10-311-455-2370	Sequence 2370, Ap
c 23	406.4	12.4	414	10	US-09-867-701-3261	Sequence 3261, Ap
24	396.6	12.1	2747	13	US-10-311-455-2369	Sequence 2369, Ap
c 25	394.4	12.0	396	10	US-09-867-701-2909	Sequence 2909, Ap
26	394.2	12.0	404	11	US-09-918-995-17414	Sequence 17414, A
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c 28	324.4	9.9	338	10	US-09-867-701-9834	Sequence 9834, Ap
c 29	306.4	9.3	330	10	US-09-867-701-3723	Sequence 3723, Ap
c 30	294.4	8.9	512	13	US-10-029-386-9769	Sequence 9769, Ap
c 31	275.8	8.4	547	13	US-10-029-386-12153	Sequence 12153, A
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c 33	271.4	8.2	278	13	US-10-029-386-23469	Sequence 23469, A
34	228.4	6.9	230	15	US-10-102-524-616	Sequence 616, App
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37	169.2	5.1	478	12	US-10-062-674-949	Sequence 949, App
38	117.6	3.6	6533	13	US-10-311-455-1681	Sequence 1681, Ap
c 39	112.6	3.4	393	15	US-10-198-846-7182	Sequence 7182, Ap
40	102.8	3.1	1764	10	US-09-938-842A-934	Sequence 934, App
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c 45	75	2.3	6533	13	US-10-311-455-1682	Sequence 1682, Ap

ALIGNMENTS

RESULT 1
US-09-981-353-172
; Sequence 172, Application US/09981353
; Patent No. US20020160382A1
; GENERAL INFORMATION:
; APPLICANT: Lasek, Amy W.
; APPLICANT: Jones, David A.
; TITLE OF INVENTION: GENES EXPRESSED IN COLON CANCER
; FILE REFERENCE: PA-0038 US
; CURRENT APPLICATION NUMBER: US/09/981,353
; CURRENT FILING DATE: 2001-10-11
; NUMBER OF SEQ ID NOS: 194
; SOFTWARE: PERL Program
; SEQ ID NO 172
; LENGTH: 3290
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No. US20020160382A1 2023119CB1
US-09-981-353-172

Query Match 100.0%; Score 3290; DB 10; Length 3290;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 3290; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Qy      241 AGCCAAAGAAACCCCAGACAACAGATGCCCATACGCAGCGTATAGCAGTAACTCCCCAGC 300
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Qy      301 TCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTTATATAATATATATTATTTATTATA 360
Db      301 TCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTTATATAATATATATTATTTATTATA 360

Qy      361 GCATTTTTGATACCTCATATTCTGTTTACACATCTTGAAAGGCGCTCAGTAGTTCTCTTA 420
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Qy       2521 CCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATTACAGTGTTAACAGAAGACTGACAA 2580
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Qy       2821 TCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAGCATGCTCTGCGTTGTTGGTTTCAC 2880
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RESULT 2
US-09-919-039-164
; Sequence 164, Application US/09919039
; Publication No. US20030108871A1
; GENERAL INFORMATION:
;  APPLICANT: Kaser, Matthew R.
;  TITLE OF INVENTION: GENES EXPRESSED IN TREATED HUMAN C3A LIVER CELL CULTURES
;  FILE REFERENCE: PA-0035 US
;  CURRENT APPLICATION NUMBER: US/09/919,039
;  CURRENT FILING DATE:  2002-09-09
;  PRIOR APPLICATION NUMBER: 60/222,113
;  PRIOR FILING DATE: 2000-07-28
;  NUMBER OF SEQ ID NOS: 401
;  SOFTWARE: PERL Program
; SEQ ID NO 164
;  LENGTH: 3290
;  TYPE: DNA
;  ORGANISM: Homo sapiens
;  FEATURE:
;  NAME/KEY: misc_feature
;  OTHER INFORMATION: Incyte ID No. US20030108871A1 2023119CB1
US-09-919-039-164

  Query Match             100.0%;  Score 3290;  DB 11;  Length 3290;
  Best Local Similarity   100.0%;  Pred. No. 0;
  Matches 3290;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

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Qy        361 GCATTTTTGATACCTCATATTCTGTTTACACATCTTGAAAGGCGCTCAGTAGTTCTCTTA 420
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Db        361 GCATTTTTGATACCTCATATTCTGTTTACACATCTTGAAAGGCGCTCAGTAGTTCTCTTA 420

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Qy        541 TTGTCTTGGCATTCTCCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGG 600
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy        661 GCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGG 720
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Db        661 GCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGG 720

Qy        721 AGATGTACAACTCGACTCAAGGGCTACTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTT 780
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Qy        781 CTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGCTCCCTATTTCTGGAACCCATTGTA 840
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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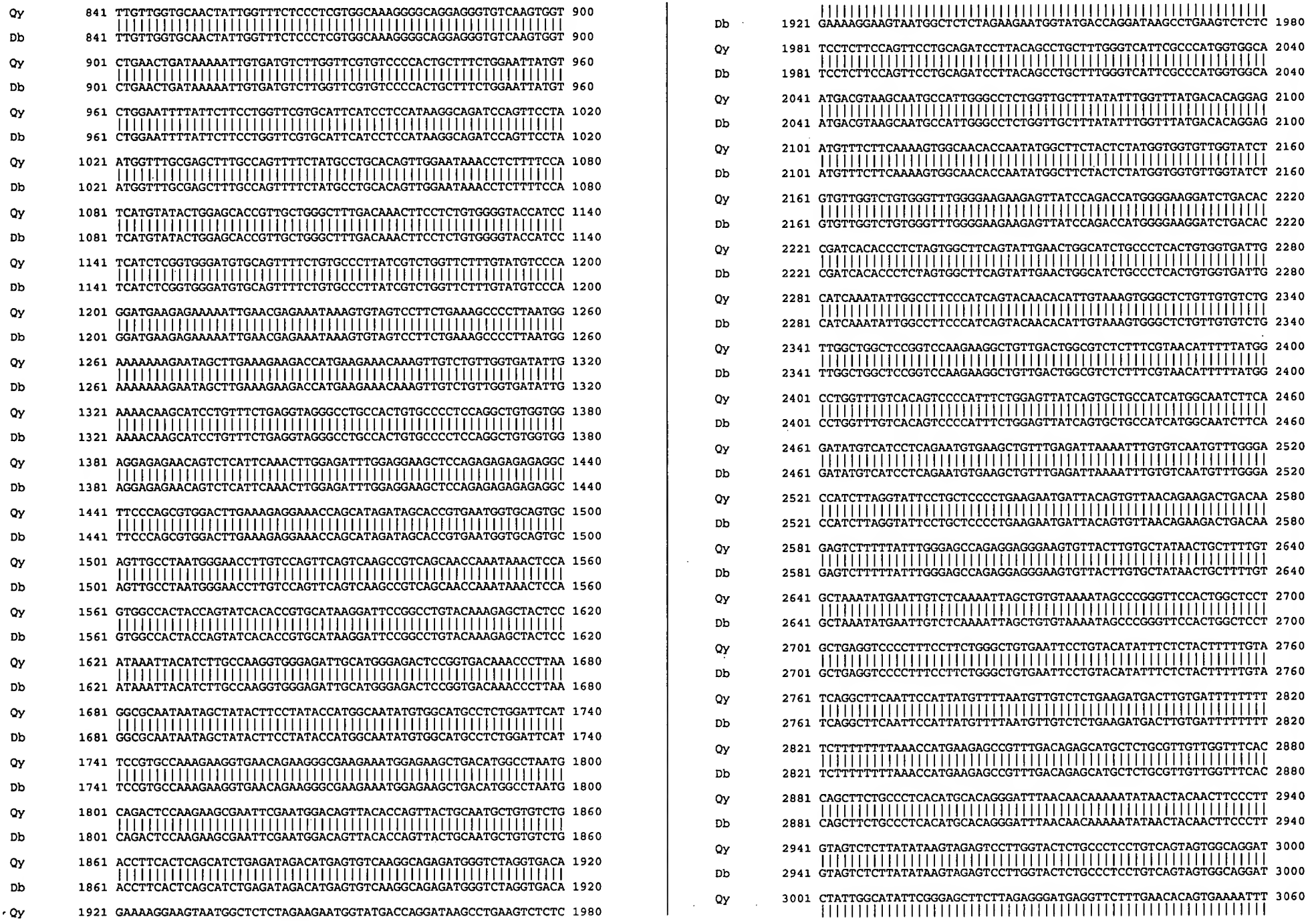
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Db 841 TTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGT 900
Qy 901 CTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGT 960
Db 901 CTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGT 960
Qy 961 CTGGAATTTTATTCTTCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTA 1020
Db 961 CTGGAATTTTATTCTTCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTA 1020
Qy 1021 ATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCA 1080
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Qy 1141 TCATCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCA 1200
Db 1141 TCATCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCA 1200
Qy 1201 GGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGG 1260
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Qy 1261 AAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTG 1320
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Db 1381 AGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGC 1440
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Qy 1801 CAGACTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTG 1860
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Qy 1861 ACCTTCACTCAGCATCTGAGATAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTGACA 1920
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Qy 2161 GTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACAC 2220
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Qy 2761 TCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTCTGAAGATGACTTGTGATTTTTTTT 2820
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Qy 2821 TCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAGCATGCTCTGCGTTGTTGGTTTCAC 2880
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RESULT 3
US-10-240-965-111
; Sequence 111, Application US/10240965
; Publication No. US20030165924A1
; GENERAL INFORMATION:
;  APPLICANT: INCYTE GENOMICS, INC.
;  APPLICANT:  SHIFFMAN, Dov
;  APPLICANT:  SOMOGYI, Roland
;  APPLICANT:  LAWN, Richard M.
;  APPLICANT:  SEILHAMER, Jeffrey J.
;  APPLICANT:  PORTER, Gordon J.
;  APPLICANT:  MIKITA, Thomas
;  APPLICANT:  TAI, Julie
;  TITLE OF INVENTION: GENES EXPRESSED IN FOAM CELL DIFFERENTIATION
;  FILE REFERENCE: PA-0025 PCT
;  CURRENT APPLICATION NUMBER: US/10/240,965
;  CURRENT FILING DATE:  2002-10-04
;  PRIOR APPLICATION NUMBER: 60/195,106
;  PRIOR FILING DATE: 2000-04-05
;  NUMBER OF SEQ ID NOS: 276
;  SOFTWARE: PERL Program
; SEQ ID NO 111
;   LENGTH: 3328
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   OTHER INFORMATION: Incyte ID No. US20030165924A1 474592.3
US-10-240-965-111

  Query Match             99.4%;  Score 3270.8;  DB 13;  Length 3328;
  Best Local Similarity   99.9%;  Pred. No. 0;
  Matches 3283;  Conservative    0;  Mismatches    2;  Indels    1;  Gaps    1;

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Qy        661 GCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGG 720
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Qy        721 AGATGTACAACTCGACTCAAGGGCTACTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTT 780
              |||||||||||||||||||||| ||||||||||||||||||||||||||||||||||||
Db        754 AGATGTACAACTCGACTCAAGGGCTGCTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTT 813

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Db        814 CTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGCTCCCTATTTCTGGAACCCATTGTA 873

Qy        841 TTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGT 900
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Db        874 TTGTTGTTGCAACTATTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGT 933

Qy        901 CTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGT 960
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        934 CTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGT 993

Qy        961 CTGGAATTTTATTCTTCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTA 1020
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        994 CTGGAATTTTATTCTTCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTA 1053

Qy       1021 ATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCA 1080
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1054 ATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCA 1113

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Db       1114 TCATGTATACTGGAGCACCGTTGCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCC 1173

Qy       1141 TCATCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCA 1200
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1174 TCATCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCA 1233

Qy       1201 GGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGG 1260
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1234 GGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGG 1293

Qy       1261 AAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTG 1320
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1294 AAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTG 1353

Qy       1321 AAAACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCACTGTGCCCCCTCCAGGCTGTGGTGG 1380
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy       1381 AGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGC 1440
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Qy      1441 TTCCCAGCGTGGACTTGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGC 1500

Db      1474 TTCCCAGCGTGGACTTGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGC 1533

Qy      1501 AGTTGCCTAATGGGAACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCA 1560

Db      1534 AGTTGCCTAATGGGAACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCA 1593

Qy      1561 GTGGCCACTACCAGTATCACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCC 1620

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Qy      1741 TCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATG 1800

Db      1774 TCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATG 1833

Qy      1801 CAGACTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTG 1860

Db      1834 CAGACTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTG 1893

Qy      1861 ACCTTCACTCAGCATCTGAGATAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTGACA 1920

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Qy      1921 GAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATGACCAGGATAAGCCTGAAGTCTCTC 1980

Db      1954 GAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATGACCAGGATAAGCCTGAAGTCTCTC 2013

Qy      1981 TCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTGGCA 2040

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Qy      2041 ATG-ACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAGGA 2099

Db      2074 ATGCACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAGGA 2133

Qy      2100 GATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATC 2159

Db      2134 GATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATC 2193

Qy      2160 TGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACA 2219

Db      2194 TGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACA 2253

Qy      2220 CCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATT 2279

Db      2254 CCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATT 2313

Qy      2280 GCATCAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGTCT 2339

Db      2314 GCATCAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGTCT 2373

Qy      2340 GTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTATG 2399

Db      2374 GTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTATG 2433

Qy      2400 GCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCAGTGCTGCCATCATGGCAATCTTC 2459

Db      2434 GCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCAGTGCTGCCATCATGGCAATCTTC 2493

Qy      2460 AGATATGTCATCCTCAGAATGTGAAGCTGTTTGAGATTAAAATTTGTGTCAATGTTTGGG 2519

Db      2494 AGATATGTCATCCTCAGAATGTGAAGCTGTTTGAGATTAAAATTTGTGTCAATGTTTGGG 2553

Qy      2520 ACCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATTACAGTGTTAACAGAAGACTGACA 2579

Db      2554 ACCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATTACAGTGTTAACAGAAGACTGACA 2613

Qy      2580 AGAGTCTTTTTATTTGGGAGCCAGAGGAGGGAAGTGTTACTTGTGCTATAACTGCTTTTG 2639

Db      2614 AGAGTCTTTTTATTTGGGAGCCAGAGGAGGGAAGTGTTACTTGTGCTATAACTGCTTTTG 2673

Qy      2640 TGCTAAATATGAATTGTCTCAAAATTAGCTGTGTAAAATAGCCCGGGTTCCACTGGCTCC 2699

Db      2674 TGCTAAATATGAATTGTCTCAAAATTAGCTGTGTAAAATAGCCCGGGTTCCACTGGCTCC 2733

Qy      2700 TGCTGAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCTGTACATATTTCTCTACTTTTTGT 2759

Db      2734 TGCTGAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCTGTACATATTTCTCTACTTTTTGT 2793

Qy      2760 ATCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTCTGAAGATGACTTGTGATTTTTTT 2819

Db      2794 ATCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTCTGAAGATGACTTGTGATTTTTTT 2853

Qy      2820 TTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAGCATGCTCTGCGTTGTTGGTTTCA 2879

Db      2854 TTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAGCATGCTCTGCGTTGTTGGTTTCA 2913

Qy      2880 CCAGCTTCTGCCCTCACATGCACAGGGATTTAACAACAAAAATATAACTACAACTTCCCT 2939

Db      2914 CCAGCTTCTGCCCTCACATGCACAGGGATTTAACAACAAAAATATAACTACAACTTCCCT 2973

Qy      2940 TGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTGCCCTCCTGTCAGTAGTGGCAGGA 2999

Db      2974 TGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTGCCCTCCTGTCAGTAGTGGCAGGA 3033

Qy      3000 TCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAGGTTCTTTGAACACAGTGAAAATT 3059

Db      3034 TCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAGGTTCTTTGAACACAGTGAAAATT 3093

Qy      3060 TAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTGTTATTGCTAAGAAGAAGTAAGAA 3119

Db      3094 TAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTGTTATTGCTAAGAAGAAGTAAGAA 3153

Qy      3120 AGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAAGATTTCTGGCAGTGTGGGATGGA 3179

Db      3154 AGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAAGATTTCTGGCAGTGTGGGATGGA 3213

Qy      3180 TGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAATGGGACAGCCTTCCATGTTCATTT 3239

Db      3214 TGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAATGGGACAGCCTTCCATGTTCATTT 3273

Qy      3240 GTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTTTAGAAAAAA 3285

Db      3274 GTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTTTAGAAAAAA 3319
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RESULT 4
US-09-954-531-137
; Sequence 137, Application US/09954531
; Patent No. US20020165180A1
; GENERAL INFORMATION:
;  APPLICANT: Weaver, Zoe
;  TITLE OF INVENTION: Process for Identifying Anti-Cancer Therapeutic Agents Using Cance
;  TITLE OF INVENTION:  Gene Sets
;  FILE REFERENCE: 689290-77
;  CURRENT APPLICATION NUMBER: US/09/954,531
;  CURRENT FILING DATE:  2002-05-02
;  PRIOR APPLICATION NUMBER: US/60/233,133
;  PRIOR FILING DATE: 2000-09-18
;  PRIOR APPLICATION NUMBER: US/60/234,009
;  PRIOR FILING DATE: 2000-09-20
;  PRIOR APPLICATION NUMBER: US/60/234,034
;  PRIOR FILING DATE: 2000-09-20
;  PRIOR APPLICATION NUMBER: US/60/234,509
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;  PRIOR FILING DATE: 2000-09-22
;  PRIOR APPLICATION NUMBER: US/60/234,567
;  PRIOR FILING DATE: 2000-09-22
;  NUMBER OF SEQ ID NOS: 1392
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 137
;   LENGTH: 3220
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-954-531-137

  Query Match             96.5%;  Score 3175.2;  DB 10;  Length 3220;
  Best Local Similarity   99.8%;  Pred. No. 0;
  Matches 3182;  Conservative    0;  Mismatches    6;  Indels    1;  Gaps    1;

Qy         97 CCGACCCGGGCCGTGTGCCCGTGGCTCCAGCCGCTGCCGCCTCGATCTCCTCGTCTCCCG 156
              ||| ||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         24 CCGGGCCGTGCCGTGTGCCCGTGGCTCCAGCCGCTGCCGCCTCGATCTCCTCGTCTCCCG 83

Qy        157 CTCCGCCCTCCCTTTTCCCTGGATGAACTTGCGTCCTTTCTCTTCTCCGCCATGGAATTC 216
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         84 CTCCGCCCTCCCTTTTCCCTGGATGAACTTGCGTCCTTTCTCTTCTCCGCCATGGAATTC 143

Qy        217 TGCTCCGTGCTTTTAGCCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATACGC 276
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        144 TGCTCCGTGCTTTTAGCCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATACGC 203

Qy        277 AGCGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTT 336
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        204 AGCGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTT 263

Qy        337 ATATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTT 396
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        264 ATATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTT 323

Qy        397 GAAAGGCGCTCAGTAGTTCTCTTACTAAACAACCACTACTCCAGAGAATGGCAACGGTGA 456
              |||||||||||||||||||||||||||||||||||||||||||||||||||| |||||||
Db        324 GAAAGGCGCTCAGTAGTTCTCTTACTAAACAACCACTACTCCAGAGAATGGCAACGCTGA 383

Qy        457 TTACCAGTACTACAGCTGCTACCGCCGCTTCTGGTCCTTTGGTGGACTACCTATGGATGC 516
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        384 TTACCAGTACTACAGCTGCTACCGCCGCTTCTGGTCCTTTGGTGGACTACCTATGGATGC 443

Qy        517 TCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCTCCGTGGGAGCCAATGATGTAG 576
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        444 TCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCTCCGTGGGAGCCAATGATGTAG 503

Qy        577 CAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAAGCAAGCCTGCATCC 636
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        504 CAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAAGCAAGCCTGCATCC 563

Qy        637 TAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCA 696
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        564 TAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCA 623

Qy        697 TCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGGGCTACTGATGGCCG 756
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        624 TCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGGGCTACTGATGGCCG 683

Qy        757 GCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGC 816
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        684 GCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGC 743

Qy        817 TCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAA 876
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        744 TCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAA 803

Qy        877 AGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGT 936
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        804 AGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGT 863

Qy        937 CCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGTTCGTGCATTCATCC 996
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

Db        864 CCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGTTCGTGCATTCATCC 923

Qy        997 TCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCA 1056
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        924 TCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCA 983

Qy       1057 CAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAGCACCGTTGCTGGGCTTTGACA 1116
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        984 CAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAGCACCGTTGCTGGGCTTTGACA 1043

Qy       1117 AACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTA 1176
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1044 AACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTA 1103

Qy       1177 TCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTA 1236
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1104 TCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTA 1163

Qy       1237 GTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAA 1296
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1164 GTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAA 1223

Qy       1297 CAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCA 1356
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1224 CAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCA 1283

Qy       1357 CTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGG 1416
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1284 CTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGG 1343

Qy       1417 AGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACTTGAAAGAGGAAACCAGCATAG 1476
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1344 AGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACTTGAAAGAGGAAACCAGCATAG 1403

Qy       1477 ATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGAACCTTGTCCAGTTCAGTCAAG 1536
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1404 ATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGAACCTTGTCCAGTTCAGTCAAG 1463

Qy       1537 CCGTCAGCAACCAAATAAACTCCAGTGGCCACTACCAGTATCACACCGTGCATAAGGATT 1596
              |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db       1464 CCGTCAGCAACCAAATAAACTCCAGTGGCCACTCCCAGTATCACACCGTGCATAAGGATT 1523

Qy       1597 CCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTTGCCAAGGTGGGAGATTGCATGG 1656
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1524 CCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTTGCCAAGGTGGGAGATTGCATGG 1583

Qy       1657 GAGACTCCGGTGACAAACCCTTAAGGCGCAATAATAGCTATACTTCCTATACCATGGCAA 1716
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1584 GAGACTCCGGTGACAAACCCTTAAGGCGCAATAATAGCTATACTTCCTATACCATGGCAA 1643

Qy       1717 TATGTGGCATGCCTCTGGATTCATTCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAA 1776
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1644 TATGTGGCATGCCTCTGGATTCATTCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAA 1703

Qy       1777 TGGAGAAGCTGACATGGCCTAATGCAGACTCCAAGAAGCGAATTCGAATGGACAGTTACA 1836
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1704 TGGAGAAGCTGACATGGCCTAATGCAGACTCCAAGAAGCGAATTCGAATGGACAGTTACA 1763

Qy       1837 CCAGTTACTGCAATGCTGTGTCTGACCTTCACTCAGCATCTGAGATAGACATGAGTGTCA 1896
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1764 CCAGTTACTGCAATGCTGTGTCTGACCTTCACTCAGCATCTGAGATAGACATGAGTGTCA 1823

Qy       1897 AGGCAGAGATGGGTCTAGGTGACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATG 1956
              |||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1824 AGGCAGCGATGGGTCTAGGTGACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATG 1883

Qy       1957 ACCAGGATAAGCCTGAAGTCTCTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCT 2016
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1884 ACCAGGATAAGCCTGAAGTCTCTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCT 1943

Qy       2017 TTGGGTCATTCGCCCATGGTGGCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTT 2076
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1944 TTGGGTCATTCGCCCATGGTGGCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTT 2003
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Qy       2077 TATATTTGGTTTATGACACAGGAGATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTC 2136
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2004 TATATTTGGTTTATGACACAGGAGATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTC 2063

Qy       2137 TACTCTATGGTGGTGTTGGTATCTGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCC 2196
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2064 TACTCTATGGTGGTGTTGGTATCTGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCC 2123

Qy       2197 AGACCATGGGGAAGGATCTGACACCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGG 2256
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2124 AGACCATGGGGAAGGATCTGACACCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGG 2183

Qy       2257 CATCTGCCCTCACTGTGGTGATTGCATCAAATATTGGCCTTCCCATCAGTACAACACATT 2316
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2184 CATCTGCCCTCACTGTGGTGATTGCATCAAATATTGGCCTTCCCATCAGTACAACACATT 2243

Qy       2317 GTAAAGTGGGCTCTGTTGTGTCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGC 2376
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2244 GTAAAGTGGGCTCTGTTGTGTCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGC 2303

Qy       2377 GTCTCTTTCGTAACATTTTTATGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCA 2436
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2304 GTCTCTTTCGTAACATTTTTATGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCA 2363

Qy       2437 GTGCTGCCATCATGGCAATCTTCAGATATGTCATCCTCAGAATGTGAAGCTGTTTGAGAT 2496
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2364 GTGCTGCCATCATGGCAATCTTCAGATATGTCATCCTCAGAATGTGAAGCTGTTTGAGAT 2423

Qy       2497 TAAAATTTGTGTCAATGTTTGGGACCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATT 2556
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2424 TAAAATTTGTGTCAATGTTTGGGACCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATT 2483

Qy       2557 ACAGTGTTAACAGAAGACTGACAAGAGTCTTTTTATTTGGGAGCCAGAGGAGGGAAGTGT 2616
             |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||
Db       2484 ACAGTGTTAACAGAAGACTGACAAGAGTCTTTTTATTTGGGAG-CAGAGGAGGGAAGTGT 2542

Qy       2617 TACTTGTGCTATAACTGCTTTTGTGCTAAATATGAATTGTCTCAAAATTAGCTGTGTAAA 2676
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2543 TACTTGTGCTATAACTGCTTTTGTGCTAAATATGAATTGTCTCAAAATTAGCTGTGTAAA 2602

Qy       2677 ATAGCCCGGGTTCCACTGGCTCCTGCTGAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCT 2736
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2603 ATAGCCCGGGTTCCACTGGCTCCTGCTGAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCT 2662

Qy       2737 GTACATATTTCTCTACTTTTTGTATCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTC 2796
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2663 GTACATATTTCTCTACTTTTTGTATCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTC 2722

Qy       2797 TGAAGATGACTTGTGATTTTTTTTTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAG 2856
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2723 TGAAGATGACTTGTGATTTTTTTTTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAG 2782

Qy       2857 CATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAAC 2916
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Db       2783 CATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAAC 2842

Qy       2917 AAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTG 2976
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2843 AAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTG 2902

Qy       2977 CCCTCCTGTCAGTAGTGGCAGGATCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAG 3036
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Db       2903 CCCTCCTGTCAGTAGTGGCAGGATCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAG 2962

Qy       3037 GTTCTTTGAACACAGTGAAAATTTAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTG 3096
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2963 GTTCTTTGAACACAGTGAAAATTTAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTG 3022

Qy       3097 TTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAA 3156
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Db       3023 TTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAA 3082

Qy       3157 GATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAAT 3216
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3083 GATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAAT 3142

Qy       3217 GGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTT 3276
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Db       3143 GGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTT 3202

Qy       3277 TAGAAAAAA 3285
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Db       3203 TAGAAAAAA 3211
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RESULT 5
US-09-954-531-356
; Sequence 356, Application US/09954531
; Patent No. US20020165180A1
; GENERAL INFORMATION:
; APPLICANT: Weaver, Zoe
; TITLE OF INVENTION: Process for Identifying Anti-Cancer Therapeutic Agents Using Cance
; TITLE OF INVENTION: Gene Sets
; FILE REFERENCE: 689290-77
; CURRENT APPLICATION NUMBER: US/09/954,531
; CURRENT FILING DATE: 2002-05-02
; PRIOR APPLICATION NUMBER: US/60/233,133
; PRIOR FILING DATE: 2000-09-18
; PRIOR APPLICATION NUMBER: US/60/234,009
; PRIOR FILING DATE: 2000-09-20
; PRIOR APPLICATION NUMBER: US/60/234,034
; PRIOR FILING DATE: 2000-09-20
; PRIOR APPLICATION NUMBER: US/60/234,509
; PRIOR FILING DATE: 2000-09-22
; PRIOR APPLICATION NUMBER: US/60/234,567
; PRIOR FILING DATE: 2000-09-22
; NUMBER OF SEQ ID NOS: 1392
; SOFTWARE: PatentIn version 3.0
; SEQ ID NO 356
; LENGTH: 3220
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-954-531-356

Query Match 96.5%; Score 3175.2; DB 10; Length 3220;
Best Local Similarity 99.8%; Pred. No. 0;
Matches 3182; Conservative 0; Mismatches 6; Indels 1; Gaps 1;

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Qy         97 CCGACCCGGGCCGTGTGCCCGTGGCTCCAGCCGCTGCCGCCTCGATCTCCTCGTCTCCCG 156
              |||  ||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         24 CCGGGCCGTGCCGTGTGCCCGTGGCTCCAGCCGCTGCCGCCTCGATCTCCTCGTCTCCCG 83

Qy        157 CTCCGCCCTCCCTTTTCCCTGGATGAACTTGCGTCCTTTCTCTTCTCCGCCATGGAATTC 216
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Db         84 CTCCGCCCTCCCTTTTCCCTGGATGAACTTGCGTCCTTTCTCTTCTCCGCCATGGAATTC 143

Qy        217 TGCTCCGTGCTTTTAGCCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATACGC 276
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        144 TGCTCCGTGCTTTTAGCCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATACGC 203

Qy        277 AGCGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTT 336
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        204 AGCGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTT 263

Qy        337 ATATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTT 396
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        264 ATATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTT 323

Qy        397 GAAAGGCGCTCAGTAGTTCTCTTACTAAACAACCACTACTCCAGAGAATGGCAACGGTGA 456
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||| |||
Db        324 GAAAGGCGCTCAGTAGTTCTCTTACTAAACAACCACTACTCCAGAGAATGGCAACGCTGA 383

Qy        457 TTACCAGTACTACAGCTGCTACCGCCGCTTCTGGTCCTTTGGTGGACTACCTATGGATGC 516
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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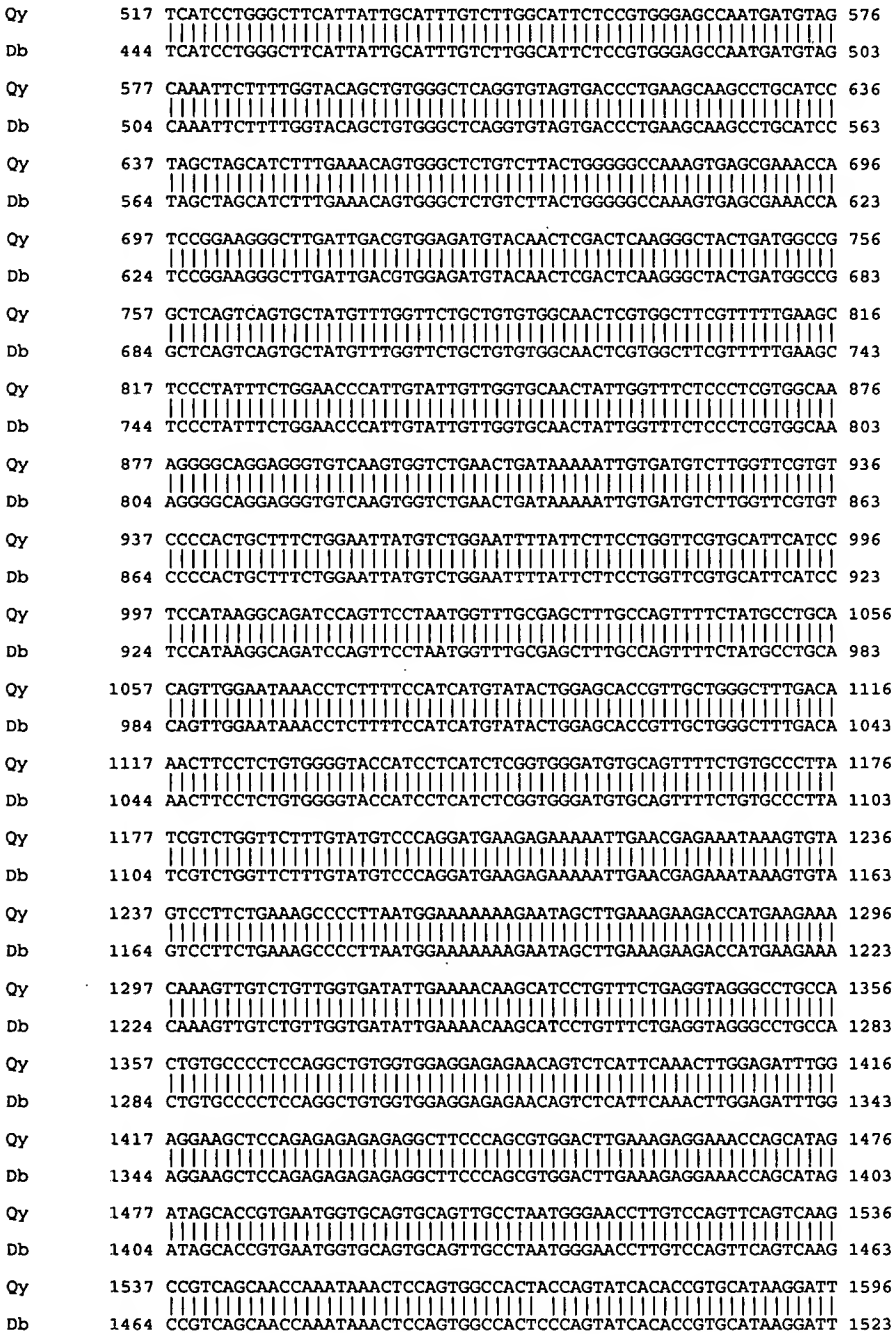
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Qy 1057 CAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAGCACCGTTGCTGGGCTTTGACA 1116
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RESULT 6
US-10-314-790-2
; Sequence 2, Application US/10314790
; Publication No. US20040002079A1
;   GENERAL INFORMATION:
;        APPLICANT: EMORY UNIVERSITY
;                   Gunn, Robert B.
;                   Timmer, Richard T.
;        TITLE OF INVENTION: SODIUM-PHOSPHATE COTRANSPORTER IN
;                            LITHIUM THERAPY FOR THE TREATMENT OF MENTAL ILLNESS
;        NUMBER OF SEQUENCES: 12
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: Kilpatrick Stockton LLP
;             STREET: Suite 2800, 1100 Peachtree Street NE
;             CITY: Atlanta
;             STATE: GA
;             COUNTRY: USA
;             ZIP: 30309-4530
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: Floppy disk
;             COMPUTER: IBM PC compatible
;             OPERATING SYSTEM: PC-DOS/MS-DOS
;             SOFTWARE: PatentIn Release #1.0, Version #1.25
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/10/314,790
;             FILING DATE: 09-Dec-2002
;             CLASSIFICATION: <Unknown>
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: US/09/380,164
;             FILING DATE: <Unknown>
;             APPLICATION NUMBER: PCT/US98/02875
;             FILING DATE: <Unknown>
;        ATTORNEY/AGENT INFORMATION:
;             NAME: Meredith, Roy D.
;             REGISTRATION NUMBER: 30,777
;             REFERENCE/DOCKET NUMBER: EMU153PCT
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: (404) 815-6500
;             TELEFAX: (404) 815-6555
;   INFORMATION FOR SEQ ID NO: 2:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 3220 base pairs
;             TYPE: nucleic acid
;             STRANDEDNESS: single
;             TOPOLOGY: linear
;        MOLECULE TYPE: mRNA
;        HYPOTHETICAL: NO
;        ANTI-SENSE: NO
;        FEATURE:
;             NAME/KEY:  hPit-1
;             LOCATION:  1..3220
;             OTHER INFORMATION:  /product= "Leukemia Virus Receptor 1"
;        PUBLICATION INFORMATION:
;             DOCUMENT NUMBER:  US 5,414,076 P
;        SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-10-314-790-2

  Query Match             96.5%;  Score 3175.2;  DB 12;  Length 3220;
  Best Local Similarity   99.8%;  Pred. No. 0;
  Matches 3182;  Conservative    0;  Mismatches    6;  Indels    1;  Gaps    1;

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                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Db        204 AGCGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTT 263

Qy        337 ATATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTT 396
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Qy        517 TCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCTCCGTGGGAGCCAATGATGTAG 576
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy        637 TAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCA 696
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
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Qy        697 TCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGGGCTACTGATGGCCG 756
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Db        684 GCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGC 743
Qy        817 TCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAA 876
Db        744 TCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAA 803
Qy        877 AGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGT 936
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Qy        997 TCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCA 1056
Db        924 TCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCA 983
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Qy       1237 GTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAA 1296
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Db       2663 GTACATATTTCTCTACTTTTTGTATCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTC 2722
Qy       2797 TGAAGATGACTTGTGATTTTTTTTTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAG 2856
Db       2723 TGAAGATGACTTGTGATTTTTTTTTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAG 2782
Qy       2857 CATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAAC 2916
Db       2783 CATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAAC 2842
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Qy       2917 AAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTG 2976
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Db       2843 AAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTG 2902

Qy       2977 CCCTCCTGTCAGTAGTGGCAGGATCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAG 3036
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2903 CCCTCCTGTCAGTAGTGGCAGGATCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAG 2962

Qy       3037 GTTCTTTGAACACAGTGAAAATTTAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTG 3096
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2963 GTTCTTTGAACACAGTGAAAATTTAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTG 3022

Qy       3097 TTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAA 3156
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Db       3023 TTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAA 3082

Qy       3157 GATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAAT 3216
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Db       3083 GATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAAT 3142

Qy       3217 GGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTT 3276
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Db       3143 GGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTT 3202

Qy       3277 TAGAAAAAA 3285
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Db       3203 TAGAAAAAA 3211
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RESULT 7
US-10-021-660-3
; Sequence 3, Application US/10021660
; Publication No. US20030152926A1
; GENERAL INFORMATION:
; APPLICANT: Murray, Richard
; APPLICANT: Glynne, Richard
; APPLICANT: Watson, Susan R.
; APPLICANT: EOS Biotechnology, Inc.
; TITLE OF INVENTION: No. US20030152926A1el Methods of Diagnosis of Angiogenesis,
; TITLE OF INVENTION: Compositions and Methods of Screening for Angiogenesis
; TITLE OF INVENTION: Modulators
; FILE REFERENCE: 018501-000710US
; CURRENT APPLICATION NUMBER: US/10/021,660
; CURRENT FILING DATE: 2001-12-06
; PRIOR APPLICATION NUMBER: US/09/784,356
; PRIOR FILING DATE: 2001-02-14
; PRIOR APPLICATION NUMBER: US 09/637,977
; PRIOR FILING DATE: 2000-08-11
; NUMBER OF SEQ ID NOS: 135
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 3
; LENGTH: 3220
; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-021-660-3

Query Match 96.5%; Score 3175.2; DB 13; Length 3220;
Best Local Similarity 99.8%; Pred. No. 0;
Matches 3182; Conservative 0; Mismatches 6; Indels 1; Gaps 1;

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Qy         97 CCGACCCGGGCCGTGTGCCCGTGGCTCCAGCCGCTGCCGCCTCGATCTCCTCGTCTCCCG 156
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Db         24 CCGGGCCGTGCCGTGTGCCCGTGGCTCCAGCCGCTGCCGCCTCGATCTCCTCGTCTCCCG 83

Qy        157 CTCCGCCCTCCCTTTTCCCTGGATGAACTTGCGTCCTTTCTCTTCTCCGCCATGGAATTC 216
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         84 CTCCGCCCTCCCTTTTCCCTGGATGAACTTGCGTCCTTTCTCTTCTCCGCCATGGAATTC 143

Qy        217 TGCTCCGTGCTTTTAGCCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATACGC 276
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        144 TGCTCCGTGCTTTTAGCCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATACGC 203

Qy        277 AGCGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTT 336
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        204 AGCGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTT 263

Qy        337 ATATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTT 396
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        264 ATATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTT 323

Qy        397 GAAAGGCGCTCAGTAGTTCTCTTACTAAACAACCACTACTCCAGAGAATGGCAACGGTGA 456
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||| |||
Db        324 GAAAGGCGCTCAGTAGTTCTCTTACTAAACAACCACTACTCCAGAGAATGGCAACGCTGA 383

Qy        457 TTACCAGTACTACAGCTGCTACCGCCGCTTCTGGTCCTTTGGTGGACTACCTATGGATGC 516
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        384 TTACCAGTACTACAGCTGCTACCGCCGCTTCTGGTCCTTTGGTGGACTACCTATGGATGC 443

Qy        517 TCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCTCCGTGGGAGCCAATGATGTAG 576
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        444 TCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCTCCGTGGGAGCCAATGATGTAG 503

Qy        577 CAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAAGCAAGCCTGCATCC 636
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        504 CAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAAGCAAGCCTGCATCC 563

Qy        637 TAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCA 696
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        564 TAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCA 623

Qy        697 TCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGGGCTACTGATGGCCG 756
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        624 TCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGGGCTACTGATGGCCG 683

Qy        757 GCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGC 816
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Db        684 GCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGC 743

Qy        817 TCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAA 876
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Db        744 TCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAA 803

Qy        877 AGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGT 936
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        804 AGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGT 863

Qy        937 CCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGTTCGTGCATTCATCC 996
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        864 CCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGTTCGTGCATTCATCC 923

Qy        997 TCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCA 1056
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        924 TCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCA 983

Qy       1057 CAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAGCACCGTTGCTGGGCTTTGACA 1116
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        984 CAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAGCACCGTTGCTGGGCTTTGACA 1043

Qy       1117 AACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTA 1176
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1044 AACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTA 1103

Qy       1177 TCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTA 1236
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1104 TCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTA 1163

Qy       1237 GTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAA 1296
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1164 GTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAA 1223

Qy       1297 CAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCA 1356
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1224 CAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCA 1283

Qy       1357 CTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGG 1416
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Db      1284 CTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGG 1343
Qy      1417 AGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACTTGAAAGAGGAAACCAGCATAG 1476
Db      1344 AGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACTTGAAAGAGGAAACCAGCATAG 1403
Qy      1477 ATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGAACCTTGTCCAGTTCAGTCAAG 1536
Db      1404 ATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGAACCTTGTCCAGTTCAGTCAAG 1463
Qy      1537 CCGTCAGCAACCAAATAAACTCCAGTGGCCACTACCAGTATCACACCGTGCATAAGGATT 1596
Db      1464 CCGTCAGCAACCAAATAAACTCCAGTGGCCACTCCCAGTATCACACCGTGCATAAGGATT 1523
Qy      1597 CCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTTGCCAAGGTGGGAGATTGCATGG 1656
Db      1524 CCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTTGCCAAGGTGGGAGATTGCATGG 1583
Qy      1657 GAGACTCCGGTGACAAACCCTTAAGGCGCAATAATAGCTATACTTCCTATACCATGGCAA 1716
Db      1584 GAGACTCCGGTGACAAACCCTTAAGGCGCAATAATAGCTATACTTCCTATACCATGGCAA 1643
Qy      1717 TATGTGGCATGCCTCTGGATTCATTCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAA 1776
Db      1644 TATGTGGCATGCCTCTGGATTCATTCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAA 1703
Qy      1777 TGGAGAAGCTGACATGGCCTAATGCAGACTCCAAGAAGCGAATTCGAATGGACAGTTACA 1836
Db      1704 TGGAGAAGCTGACATGGCCTAATGCAGACTCCAAGAAGCGAATTCGAATGGACAGTTACA 1763
Qy      1837 CCAGTTACTGCAATGCTGTGTCTGACCTTCACTCAGCATCTGAGATAGACATGAGTGTCA 1896
Db      1764 CCAGTTACTGCAATGCTGTGTCTGACCTTCACTCAGCATCTGAGATAGACATGAGTGTCA 1823
Qy      1897 AGGCAGAGATGGGTCTAGGTGACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATG 1956
Db      1824 AGGCAGCGATGGGTCTAGGTGACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATG 1883
Qy      1957 ACCAGGATAAGCCTGAAGTCTCTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCT 2016
Db      1884 ACCAGGATAAGCCTGAAGTCTCTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCT 1943
Qy      2017 TTGGGTCATTCGCCCATGGTGGCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTT 2076
Db      1944 TTGGGTCATTCGCCCATGGTGGCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTT 2003
Qy      2077 TATATTTGGTTTATGACACAGGAGATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTC 2136
Db      2004 TATATTTGGTTTATGACACAGGAGATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTC 2063
Qy      2137 TACTCTATGGTGGTGTTGGTATCTGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCC 2196
Db      2064 TACTCTATGGTGGTGTTGGTATCTGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCC 2123
Qy      2197 AGACCATGGGGAAGGATCTGACACCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGG 2256
Db      2124 AGACCATGGGGAAGGATCTGACACCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGG 2183
Qy      2257 CATCTGCCCTCACTGTGGTGATTGCATCAAATATTGGCCTTCCCATCAGTACAACACATT 2316
Db      2184 CATCTGCCCTCACTGTGGTGATTGCATCAAATATTGGCCTTCCCATCAGTACAACACATT 2243
Qy      2317 GTAAAGTGGGCTCTGTTGTGTCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGC 2376
Db      2244 GTAAAGTGGGCTCTGTTGTGTCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGC 2303
Qy      2377 GTCTCTTTCGTAACATTTTTATGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCA 2436
Db      2304 GTCTCTTTCGTAACATTTTTATGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCA 2363
Qy      2437 GTGCTGCCATCATGGCAATCTTCAGATATGTCATCCTCAGAATGTGAAGCTGTTTGAGAT 2496
Db      2364 GTGCTGCCATCATGGCAATCTTCAGATATGTCATCCTCAGAATGTGAAGCTGTTTGAGAT 2423
Qy      2497 TAAAATTTGTGTCAATGTTTGGGACCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATT 2556
Db      2424 TAAAATTTGTGTCAATGTTTGGGACCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATT 2483
Qy      2557 ACAGTGTTAACAGAAGACTGACAAGAGTCTTTTTATTTGGGAGCCAGAGGAGGGAAGTGT 2616
Db      2484 ACAGTGTTAACAGAAGACTGACAAGAGTCTTTTTATTTGGGAG-CAGAGGAGGGAAGTGT 2542
Qy      2617 TACTTGTGCTATAACTGCTTTTGTGCTAAATATGAATTGTCTCAAAATTAGCTGTGTAAA 2676
Db      2543 TACTTGTGCTATAACTGCTTTTGTGCTAAATATGAATTGTCTCAAAATTAGCTGTGTAAA 2602
Qy      2677 ATAGCCCGGGTTCCACTGGCTCCTGCTGAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCT 2736
Db      2603 ATAGCCCGGGTTCCACTGGCTCCTGCTGAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCT 2662
Qy      2737 GTACATATTTCTCTACTTTTTGTATCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTC 2796
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Qy      2797 TGAAGATGACTTGTGATTTTTTTTTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAG 2856
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Qy      2857 CATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAAC 2916
Db      2783 CATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAAC 2842
Qy      2917 AAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTG 2976
Db      2843 AAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTG 2902
Qy      2977 CCCTCCTGTCAGTAGTGGCAGGATCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAG 3036
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Qy      3037 GTTCTTTGAACACAGTGAAAATTTAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTG 3096
Db      2963 GTTCTTTGAACACAGTGAAAATTTAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTG 3022
Qy      3097 TTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAA 3156
Db      3023 TTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAA 3082
Qy      3157 GATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAAT 3216
Db      3083 GATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAAT 3142
Qy      3217 GGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTT 3276
Db      3143 GGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTT 3202
Qy      3277 TAGAAAAAA 3285
Db      3203 TAGAAAAAA 3211
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RESULT 8
US-09-873-367C-694
; Sequence 694, Application US/09873367C
; Publication No. US20030165839A1
; GENERAL INFORMATION:
; APPLICANT: Young, Paul
; APPLICANT: Soppet, Daniel
; APPLICANT: Endress, Gregory
; APPLICANT: Augustus, Meena
; APPLICANT: Ebner, Reinhard
; APPLICANT: Carter, Kenneth
; TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using
; TITLE OF INVENTION: Signature Gene Sets
; FILE REFERENCE: 689290-64

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;  CURRENT APPLICATION NUMBER: US/09/873,367C
;  CURRENT FILING DATE:  2003-04-29
;  PRIOR APPLICATION NUMBER: U.S. 60/236,891
;  PRIOR FILING DATE: 2000-09-29
;  PRIOR APPLICATION NUMBER: U.S. 60/236,842
;  PRIOR FILING DATE: 2000-09-29
;  PRIOR APPLICATION NUMBER: U.S. 60/244,867
;  PRIOR FILING DATE: 2000-11-01
;  PRIOR APPLICATION NUMBER: U.S. 60/245,084
;  PRIOR FILING DATE: 2000-11-01
;  NUMBER OF SEQ ID NOS: 1067
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 694
;   LENGTH: 3220
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-873-367C-694

  Query Match             96.5%;  Score 3175.2;  DB 13;  Length 3220;
  Best Local Similarity   99.8%;  Pred. No. 0;
  Matches 3182;  Conservative    0;  Mismatches    6;  Indels    1;  Gaps    1;

Qy         97 CCGACCCGGGCCGTGTGCCCGTGGCTCCAGCCGCTGCCGCCTCGATCTCCTCGTCTCCCG 156
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Db         24 CCGGGCCGTGCCGTGTGCCCGTGGCTCCAGCCGCTGCCGCCTCGATCTCCTCGTCTCCCG 83

Qy        157 CTCCGCCCTCCCTTTTCCCTGGATGAACTTGCGTCCTTTCTCTTCTCCGCCATGGAATTC 216
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         84 CTCCGCCCTCCCTTTTCCCTGGATGAACTTGCGTCCTTTCTCTTCTCCGCCATGGAATTC 143

Qy        217 TGCTCCGTGCTTTTAGCCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATACGC 276
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        144 TGCTCCGTGCTTTTAGCCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATACGC 203

Qy        277 AGCGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTT 336
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Db        204 AGCGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTT 263

Qy        337 ATATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTT 396
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        264 ATATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTT 323

Qy        397 GAAAGGCGCTCAGTAGTTCTCTTACTAAACAACCACTACTCCAGAGAATGGCAACGGTGA 456
              |||||||||||||||||||||||||||||||||||||||||||||||||||||| |||
Db        324 GAAAGGCGCTCAGTAGTTCTCTTACTAAACAACCACTACTCCAGAGAATGGCAACGCTGA 383

Qy        457 TTACCAGTACTACAGCTGCTACCGCCGCTTCTGGTCCTTTGGTGGACTACCTATGGATGC 516
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        384 TTACCAGTACTACAGCTGCTACCGCCGCTTCTGGTCCTTTGGTGGACTACCTATGGATGC 443

Qy        517 TCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCTCCGTGGGAGCCAATGATGTAG 576
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        444 TCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCTCCGTGGGAGCCAATGATGTAG 503

Qy        577 CAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAAGCAAGCCTGCATCC 636
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        504 CAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAAGCAAGCCTGCATCC 563

Qy        637 TAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCA 696
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        564 TAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCA 623

Qy        697 TCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGGGCTACTGATGGCCG 756
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        624 TCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGGGCTACTGATGGCCG 683

Qy        757 GCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGC 816
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Db        684 GCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGC 743

Qy        817 TCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAA 876
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        744 TCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGGCAA 803

Qy        877 AGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGT 936
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Db        804 AGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGATGTCTTGGTTCGTGT 863

Qy        937 CCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGTTCGTGCATTCATCC 996
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Db        864 CCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGTTCGTGCATTCATCC 923

Qy        997 TCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCA 1056
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Db        924 TCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCA 983

Qy       1057 CAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAGCACCGTTGCTGGGCTTTGACA 1116
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Db        984 CAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAGCACCGTTGCTGGGCTTTGACA 1043

Qy       1117 AACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTA 1176
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Db       1044 AACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTA 1103

Qy       1177 TCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTA 1236
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1104 TCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTA 1163

Qy       1237 GTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAA 1296
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1164 GTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAA 1223

Qy       1297 CAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCA 1356
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1224 CAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCA 1283

Qy       1357 CTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGG 1416
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1284 CTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGG 1343

Qy       1417 AGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACTTGAAAGAGGAAACCAGCATAG 1476
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1344 AGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACTTGAAAGAGGAAACCAGCATAG 1403

Qy       1477 ATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGAACCTTGTCCAGTTCAGTCAAG 1536
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1404 ATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGAACCTTGTCCAGTTCAGTCAAG 1463

Qy       1537 CCGTCAGCAACCAAATAAACTCCAGTGGCCACTACCAGTATCACACCGTGCATAAGGATT 1596
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db       1464 CCGTCAGCAACCAAATAAACTCCAGTGGCCACTCCCAGTATCACACCGTGCATAAGGATT 1523

Qy       1597 CCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTTGCCAAGGTGGGAGATTGCATGG 1656
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1524 CCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTTGCCAAGGTGGGAGATTGCATGG 1583

Qy       1657 GAGACTCCGGTGACAAACCCTTAAGGCGCAATAATAGCTATACTTCCTATACCATGGCAA 1716
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1584 GAGACTCCGGTGACAAACCCTTAAGGCGCAATAATAGCTATACTTCCTATACCATGGCAA 1643

Qy       1717 TATGTGGCATGCCTCTGGATTCATTCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAA 1776
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1644 TATGTGGCATGCCTCTGGATTCATTCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAA 1703

Qy       1777 TGGAGAAGCTGACATGGCCTAATGCAGACTCCAAGAAGCGAATTCGAATGGACAGTTACA 1836
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1704 TGGAGAAGCTGACATGGCCTAATGCAGACTCCAAGAAGCGAATTCGAATGGACAGTTACA 1763

Qy       1837 CCAGTTACTGCAATGCTGTGTCTGACCTTCACTCAGCATCTGAGATAGACATGAGTGTCA 1896
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1764 CCAGTTACTGCAATGCTGTGTCTGACCTTCACTCAGCATCTGAGATAGACATGAGTGTCA 1823

Qy       1897 AGGCAGAGATGGGTCTAGGTGACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATG 1956
              |||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1824 AGGCAGCGATGGGTCTAGGTGACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATG 1883
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Qy       1957 ACCAGGATAAGCCTGAAGTCTCTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCT 2016
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1884 ACCAGGATAAGCCTGAAGTCTCTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCT 1943

Qy       2017 TTGGGTCATTCGCCCATGGTGGCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTT 2076
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1944 TTGGGTCATTCGCCCATGGTGGCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTT 2003

Qy       2077 TATATTTGGTTTATGACACAGGAGATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTC 2136
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2004 TATATTTGGTTTATGACACAGGAGATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTC 2063

Qy       2137 TACTCTATGGTGGTGTTGGTATCTGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCC 2196
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2064 TACTCTATGGTGGTGTTGGTATCTGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCC 2123

Qy       2197 AGACCATGGGGAAGGATCTGACACCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGG 2256
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2124 AGACCATGGGGAAGGATCTGACACCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGG 2183

Qy       2257 CATCTGCCCTCACTGTGGTGATTGCATCAAATATTGGCCTTCCCATCAGTACAACACATT 2316
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2184 CATCTGCCCTCACTGTGGTGATTGCATCAAATATTGGCCTTCCCATCAGTACAACACATT 2243

Qy       2317 GTAAAGTGGGCTCTGTTGTGTCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGC 2376
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2244 GTAAAGTGGGCTCTGTTGTGTCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGC 2303

Qy       2377 GTCTCTTTCGTAACATTTTTATGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCA 2436
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2304 GTCTCTTTCGTAACATTTTTATGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCA 2363

Qy       2437 GTGCTGCCATCATGGCAATCTTCAGATATGTCATCCTCAGAATGTGAAGCTGTTTGAGAT 2496
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2364 GTGCTGCCATCATGGCAATCTTCAGATATGTCATCCTCAGAATGTGAAGCTGTTTGAGAT 2423

Qy       2497 TAAAATTTGTGTCAATGTTTGGGACCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATT 2556
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2424 TAAAATTTGTGTCAATGTTTGGGACCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATT 2483

Qy       2557 ACAGTGTTAACAGAAGACTGACAAGAGTCTTTTTATTTGGGAGCCAGAGGAGGGAAGTGT 2616
             |||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||
Db       2484 ACAGTGTTAACAGAAGACTGACAAGAGTCTTTTTATTTGGGAG-CAGAGGAGGGAAGTGT 2542

Qy       2617 TACTTGTGCTATAACTGCTTTTGTGCTAAATATGAATTGTCTCAAAATTAGCTGTGTAAA 2676
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2543 TACTTGTGCTATAACTGCTTTTGTGCTAAATATGAATTGTCTCAAAATTAGCTGTGTAAA 2602

Qy       2677 ATAGCCCGGGTTCCACTGGCTCCTGCTGAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCT 2736
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2603 ATAGCCCGGGTTCCACTGGCTCCTGCTGAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCT 2662

Qy       2737 GTACATATTTCTCTACTTTTTGTATCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTC 2796
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2663 GTACATATTTCTCTACTTTTTGTATCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTC 2722

Qy       2797 TGAAGATGACTTGTGATTTTTTTTTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAG 2856
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2723 TGAAGATGACTTGTGATTTTTTTTTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAG 2782

Qy       2857 CATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAAC 2916
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2783 CATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAAC 2842

Qy       2917 AAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTG 2976
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2843 AAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTG 2902

Qy       2977 CCCTCCTGTCAGTAGTGGCAGGATCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAG 3036
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2903 CCCTCCTGTCAGTAGTGGCAGGATCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAG 2962

Qy       3037 GTTCTTTGAACACAGTGAAAATTTAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTG 3096
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2963 GTTCTTTGAACACAGTGAAAATTTAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTG 3022

Qy       3097 TTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAA 3156
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3023 TTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAA 3082

Qy       3157 GATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAAT 3216
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3083 GATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAAT 3142

Qy       3217 GGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTT 3276
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3143 GGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTT 3202

Qy       3277 TAGAAAAAA 3285
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Db       3203 TAGAAAAAA 3211
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RESULT 9
US-10-024-298A-124
; Sequence 124, Application US/10024298A
; Publication No. US20030143540A1
; GENERAL INFORMATION:
; APPLICANT: ASAHI KASEI KABUSHIKI KAISHA
; APPLICANT: Akio MATSUDA
; APPLICANT: Goichi HONDA
; APPLICANT: Shuji MURAMATSU
; APPLICANT: Yukiko NAGANO
; TITLE OF INVENTION: NF-K B Activating Gene
; FILE REFERENCE: 1254-0191P
; CURRENT APPLICATION NUMBER: US/10/024,298A
; CURRENT FILING DATE: 2003-04-08
; PRIOR APPLICATION NUMBER: 60/314,385
; PRIOR FILING DATE: 2001-08-24
; PRIOR APPLICATION NUMBER: 60/278,641
; PRIOR FILING DATE: 2001-03-26
; PRIOR APPLICATION NUMBER: 60/258,315
; PRIOR FILING DATE: 2000-12-28
; PRIOR APPLICATION NUMBER: JP254018/2001
; PRIOR FILING DATE: 2001-08-24
; PRIOR APPLICATION NUMBER: JP0088912/2001
; PRIOR FILING DATE: 2001-03-26
; PRIOR APPLICATION NUMBER: JP402288/2000
; PRIOR FILING DATE: 2000-12-28
; NUMBER OF SEQ ID NOS: 182
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 124
; LENGTH: 2916
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: CDS
; LOCATION: (81)..(2117)
US-10-024-298A-124

Query Match 88.5%; Score 2911.2; DB 13; Length 2916;
Best Local Similarity 99.9%; Pred. No. 0;
Matches 2913; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

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Qy        364 TTTTTGATACCTCATATTCTGTTTACACATCTTGAAAGGCGCTCAGTAGTTCTCTTACTA 423
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          1 TTTTTGATACCTCATATTCTGTTTACACATCTTGAAAGGCGCTCAGTAGTTCTCTTACTA 60

Qy        424 AACAACCACTACTCCAGAGAATGGCAACGGTGATTACCAGTACTACAGCTGCTACCGCCG 483
              |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||
Db         61 AACAACCACTACTCCAGAGAATGGCAACGCTGATTACCAGTACTACAGCTGCTACCGCCG 120

Qy        484 CTTCTGGTCCTTTGGTGGACTACCTATGGATGCTCATCCTGGGCTTCATTATTGCATTTG 543
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 CTTCTGGTCCTTTGGTGGACTACCTATGGATGCTCATCCTGGGCTTCATTATTGCATTTG 180
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Qy       544 TCTTGGCATTCTCCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCT 603
Db       181 TCTTGGCATTCTCCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCT 240

Qy       604 CAGGTGTAGTGACCCTGAAGCAAGCCTGCATCCTAGCTAGCATCTTTGAAACAGTGGGCT 663
Db       241 CAGGTGTAGTGACCCTGAAGCAAGCCTGCATCCTAGCTAGCATCTTTGAAACAGTGGGCT 300

Qy       664 CTGTCTTACTGGGGGCCAAAGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGGAGA 723
Db       301 CTGTCTTACTGGGGGCCAAAGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGGAGA 360

Qy       724 TGTACAACTCGACTCAAGGGCTACTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTTCTG 783
Db       361 TGTACAACTCGACTCAAGGGCTGCTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTTCTG 420

Qy       784 CTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGCTCCCTATTTCTGGAACCCATTGTATTG 843
Db       421 CTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGCTCCCTATTTCTGGAACCCATTGTATTG 480

Qy       844 TTGGTGCAACTATTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTG 903
Db       481 TTGGTGCAACTATTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTG 540

Qy       904 AACTGATAAAAATTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGTCTG 963
Db       541 AACTGATAAAAATTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGTCTG 600

Qy       964 GAATTTTATTCTTCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATG 1023
Db       601 GAATTTTATTCTTCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATG 660

Qy      1024 GTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCATCA 1083
Db       661 GTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCATCA 720

Qy      1084 TGTATACTGGAGCACCGTTGCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCA 1143
Db       721 TGTATACTGGAGCACCGTTGCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCA 780

Qy      1144 TCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGA 1203
Db       781 TCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGA 840

Qy      1204 TGAAGAGAAAAATTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAA 1263
Db       841 TGAAGAGAAAAATTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAA 900

Qy      1264 AAAAGAATAGCTTGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAA 1323
Db       901 AAAAGAATAGCTTGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAA 960

Qy      1324 ACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGG 1383
Db       961 ACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGG 1020

Qy      1384 AGAGAACAGTCTCATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGCTTC 1443
Db       1021 AGAGAACAGTCTCATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGCTTC 1080

Qy      1444 CCAGCGTGGACTTGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGCAGT 1503
Db       1081 CCAGCGTGGACTTGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGCAGT 1140

Qy      1504 TGCCTAATGGGAACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTG 1563
Db       1141 TGCCTAATGGGAACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTG 1200

Qy      1564 GCCACTACCAGTATCACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCCATA 1623
Db       1201 GCCACTACCAGTATCACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCCATA 1260

Qy      1624 AATTACATCTTGCCAAGGTGGGAGATTGCATGGGAGACTCCGGTGACAAACCCTTAAGGC 1683

Db       1261 AATTACATCTTGCCAAGGTGGGAGATTGCATGGGAGACTCCGGTGACAAACCCTTAAGGC 1320

Qy      1684 GCAATAATAGCTATACTTCCTATACCATGGCAATATGTGGCATGCCTCTGGATTCATTCC 1743
Db       1321 GCAATAATAGCTATACTTCCTATACCATGGCAATATGTGGCATGCCTCTGGATTCATTCC 1380

Qy      1744 GTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATGCAG 1803
Db       1381 GTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATGCAG 1440

Qy      1804 ACTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACC 1863
Db       1441 ACTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACC 1500

Qy      1864 TTCACTCAGCATCTGAGATAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTGACAGAA 1923
Db       1501 TTCACTCAGCATCTGAGATAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTGACAGAA 1560

Qy      1924 AAGGAAGTAATGGCTCTCTAGAAGAATGGTATGACCAGGATAAGCCTGAAGTCTCTCTCC 1983
Db       1561 AAGGAAGTAATGGCTCTCTAGAAGAATGGTATGACCAGGATAAGCCTGAAGTCTCTCTCC 1620

Qy      1984 TCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTGGCAATG 2043
Db       1621 TCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTGGCAATG 1680

Qy      2044 ACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAGGAGATG 2103
Db       1681 ACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAGGAGATG 1740

Qy      2104 TTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATCTGTG 2163
Db       1741 TTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATCTGTG 1800

Qy      2164 TTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACACCGA 2223
Db       1801 TTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACACCGA 1860

Qy      2224 TCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATTGCAT 2283
Db       1861 TCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATTGCAT 1920

Qy      2284 CAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGTCTGTTG 2343
Db       1921 CAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGTCTGTTG 1980

Qy      2344 GCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTATGGCCT 2403
Db       1981 GCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTATGGCCT 2040

Qy      2404 GGTTTGTCACAGTCCCCATTTCTGGAGTTATCAGTGCTGCCATCATGGCAATCTTCAGAT 2463
Db       2041 GGTTTGTCACAGTCCCTATTTCTGGAGTTATCAGTGCTGCCATCATGGCAATCTTCAGAT 2100

Qy      2464 ATGTCATCCTCAGAATGTGAAGCTGTTTGAGATTAAAATTTGTGTCAATGTTTGGGACCA 2523
Db       2101 ATGTCATCCTCAGAATGTGAAGCTGTTTGAGATTAAAATTTGTGTCAATGTTTGGGACCA 2160

Qy      2524 TCTTAGGTATTCCTGCTCCCCTGAAGAATGATTACAGTGTTAACAGAAGACTGACAAGAG 2583
Db       2161 TCTTAGGTATTCCTGCTCCCCTGAAGAATGATTACAGTGTTAACAGAAGACTGACAAGAG 2220

Qy      2584 TCTTTTTATTTGGGAGCCAGAGGAGGGAAGTGTTACTTGTGCTATAACTGCTTTTGTGCT 2643
Db       2221 TCTTTTTATTTGGGAGCCAGAGGAGGGAAGTGTTACTTGTGCTATAACTGCTTTTGTGCT 2280

Qy      2644 AAATATGAATTGTCTCAAAATTAGCTGTGTAAAATAGCCCGGGTTCCACTGGCTCCTGCT 2703
Db       2281 AAATATGAATTGTCTCAAAATTAGCTGTGTAAAATAGCCCGGGTTCCACTGGCTCCTGCT 2340

Qy      2704 GAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCTGTACATATTTCTCTACTTTTTGTATCA 2763
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Db       2341 GAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCTGTACATATTTCTCTACTTTTTGTATCA 2400

Qy       2764 GGCTTCAATTCCATTATGTTTTAATGTTGTCTCTGAAGATGACTTGTGATTTTTTTTTCT 2823
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2401 GGCTTCAATTCCATTATGTTTTAATGTTGTCTCTGAAGATGACTTGTGATTTTTTTTTCT 2460

Qy       2824 TTTTTTTAAACCATGAAGAGCCGTTTGACAGAGCATGCTCTGCGTTGTTGGTTTCACCAG 2883
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2461 TTTTTTTAAACCATGAAGAGCCGTTTGACAGAGCATGCTCTGCGTTGTTGGTTTCACCAG 2520

Qy       2884 CTTCTGCCCTCACATGCACAGGGATTTAACAACAAAAATATAACTACAACTTCCCTTGTA 2943
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2521 CTTCTGCCCTCACATGCACAGGGATTTAACAACAAAAATATAACTACAACTTCCCTTGTA 2580

Qy       2944 GTCTCTTATATAAGTAGAGTCCTTGGTACTCTGCCCTCCTGTCAGTAGTGGCAGGATCTA 3003
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2581 GTCTCTTATATAAGTAGAGTCCTTGGTACTCTGCCCTCCTGTCAGTAGTGGCAGGATCTA 2640

Qy       3004 TTGGCATATTCGGGAGCTTCTTAGAGGGATGAGGTTCTTTGAACACAGTGAAAATTTAAA 3063
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2641 TTGGCATATTCGGGAGCTTCTTAGAGGGATGAGGTTCTTTGAACACAGTGAAAATTTAAA 2700

Qy       3064 TTAGTAACTTTTTTGCAAGCAGTTTATTGACTGTTATTGCTAAGAAGAAGTAAGAAAGAA 3123
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2701 TTAGTAACTTTTTTGCAAGCAGTTTATTGACTGTTATTGCTAAGAAGAAGTAAGAAAGAA 2760

Qy       3124 AAAGCCTGTTGGCAATCTTGGTTATTTCTTTAAGATTTCTGGCAGTGTGGGATGGATGAA 3183
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2761 AAAGCCTGTTGGCAATCTTGGTTATTTCTTTAAGATTTCTGGCAGTGTGGGATGGATGAA 2820

Qy       3184 TGAAGTGGAATGTGAACTTTGGGCAAGTTAAATGGGACAGCCTTCCATGTTCATTTGTCT 3243
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2821 TGAAGTGGAATGTGAACTTTGGGCAAGTTAAATGGGACAGCCTTCCATGTTCATTTGTCT 2880

Qy       3244 ACCTCTTAACTGAATAAAAAAGCCTACAGTTTTTAG 3279
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Db       2881 ACCTCTTAACTGAATAAAAAAGCCTACAGTTTTTAG 2916


RESULT 10
US-10-042-211A-124
; Sequence 124, Application US/10042211A
; Publication No. US20030170719A1
; GENERAL INFORMATION:
;  APPLICANT: MATSUDA, Akio et al.
;  TITLE OF INVENTION: NFKB Activating Gene
;  FILE REFERENCE: 1254-0192P
;  CURRENT APPLICATION NUMBER: US/10/042,211A
;  CURRENT FILING DATE:  2002-01-11
;  PRIOR APPLICATION NUMBER: JP 2000-402288
;  PRIOR FILING DATE: 2000-12-28
;  PRIOR APPLICATION NUMBER: JP 2001-088912
;  PRIOR FILING DATE: 2001-03-26
;  PRIOR APPLICATION NUMBER: JP 2001-254018
;  PRIOR FILING DATE: 2001-08-24
;  PRIOR APPLICATION NUMBER: US 60/258,315
;  PRIOR FILING DATE: 2000-12-28
;  PRIOR APPLICATION NUMBER: US 60/278,640
;  PRIOR FILING DATE: 2001-03-26
;  PRIOR APPLICATION NUMBER: US 60/314,385
;  PRIOR FILING DATE: 2001-08-24
;  NUMBER OF SEQ ID NOS: 182
;  SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 124
;   LENGTH: 2916
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: CDS
;   LOCATION: (81)..(2117)
US-10-042-211A-124

  Query Match             88.5%;  Score 2911.2;  DB 13;  Length 2916;
  Best Local Similarity   99.9%;  Pred. No. 0;
  Matches 2913;  Conservative    0;  Mismatches    3;  Indels    0;  Gaps    0;

Qy        364 TTTTTGATACCTCATATTCTGTTTACACATCTTGAAAGGCGCTCAGTAGTTCTCTTACTA 423
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Db          1 TTTTTGATACCTCATATTCTGTTTACACATCTTGAAAGGCGCTCAGTAGTTCTCTTACTA 60

Qy        424 AACAACCACTACTCCAGAGAATGGCAACGGTGATTACCAGTACTACAGCTGCTACCGCCG 483
              ||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||
Db         61 AACAACCACTACTCCAGAGAATGGCAACGCTGATTACCAGTACTACAGCTGCTACCGCCG 120

Qy        484 CTTCTGGTCCTTTGGTGGACTACCTATGGATGCTCATCCTGGGCTTCATTATTGCATTTG 543
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 CTTCTGGTCCTTTGGTGGACTACCTATGGATGCTCATCCTGGGCTTCATTATTGCATTTG 180

Qy        544 TCTTGGCATTCTCCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCT 603
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 TCTTGGCATTCTCCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCT 240

Qy        604 CAGGTGTAGTGACCCTGAAGCAAGCCTGCATCCTAGCTAGCATCTTTGAAACAGTGGGCT 663
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 CAGGTGTAGTGACCCTGAAGCAAGCCTGCATCCTAGCTAGCATCTTTGAAACAGTGGGCT 300

Qy        664 CTGTCTTACTGGGGGCCAAAGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGGAGA 723
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Db        301 CTGTCTTACTGGGGGCCAAAGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGGAGA 360

Qy        724 TGTACAACTCGACTCAAGGGCTACTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTTCTG 783
              ||||||||||||||||||| ||||||||||||||||||||||||||||||||||||||||||
Db        361 TGTACAACTCGACTCAAGGGCTGCTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTTCTG 420

Qy        784 CTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGCTCCCTATTTCTGGAACCCATTGTATTG 843
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        421 CTGTGTGGCAACTCGTGGCTTCGTTTTTGAAGCTCCCTATTTCTGGAACCCATTGTATTG 480

Qy        844 TTGGTGCAACTATTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTG 903
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Db        481 TTGGTGCAACTATTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTG 540

Qy        904 AACTGATAAAAATTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGTCTG 963
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Db        541 AACTGATAAAAATTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGTCTG 600

Qy        964 GAATTTTATTCTTCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATG 1023
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Db        601 GAATTTTATTCTTCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATG 660

Qy       1024 GTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCATCA 1083
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Db        661 GTTTGCGAGCTTTGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCATCA 720

Qy       1084 TGTATACTGGAGCACCGTTGCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCA 1143
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Db        721 TGTATACTGGAGCACCGTTGCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCA 780

Qy       1144 TCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGA 1203
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Db        781 TCTCGGTGGGATGTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGA 840

Qy       1204 TGAAGAGAAAAATTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAA 1263
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Qy       1264 AAAAGAATAGCTTGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAA 1323
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Db        901 AAAAGAATAGCTTGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAA 960

Qy       1324 ACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGG 1383
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Db        961 ACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGG 1020

Qy       1384 AGAGAACAGTCTCATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGCTTC 1443
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Db       1021 AGAGAACAGTCTCATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGCTTC 1080
Qy       1444 CCAGCGTGGACTTGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGCAGT 1503
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Db       1081 CCAGCGTGGACTTGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGCAGT 1140
Qy       1504 TGCCTAATGGGAACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTG 1563
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Db       1141 TGCCTAATGGGAACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTG 1200
Qy       1564 GCCACTACCAGTATCACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCCATA 1623
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Db       1201 GCCACTACCAGTATCACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCCATA 1260
Qy       1624 AATTACATCTTGCCAAGGTGGGAGATTGCATGGGAGACTCCGGTGACAAACCCTTAAGGC 1683
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Db       1261 AATTACATCTTGCCAAGGTGGGAGATTGCATGGGAGACTCCGGTGACAAACCCTTAAGGC 1320
Qy       1684 GCAATAATAGCTATACTTCCTATACCATGGCAATATGTGGCATGCCTCTGGATTCATTCC 1743
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Db       1321 GCAATAATAGCTATACTTCCTATACCATGGCAATATGTGGCATGCCTCTGGATTCATTCC 1380
Qy       1744 GTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATGCAG 1803
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Db       1381 GTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATGCAG 1440
Qy       1804 ACTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACC 1863
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Db       1441 ACTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACC 1500
Qy       1864 TTCACTCAGCATCTGAGATAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTGACAGAA 1923
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Db       1501 TTCACTCAGCATCTGAGATAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTGACAGAA 1560
Qy       1924 AAGGAAGTAATGGCTCTCTAGAAGAATGGTATGACCAGGATAAGCCTGAAGTCTCTCTCC 1983
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Db       1561 AAGGAAGTAATGGCTCTCTAGAAGAATGGTATGACCAGGATAAGCCTGAAGTCTCTCTCC 1620
Qy       1984 TCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTGGCAATG 2043
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Db       1621 TCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTGGCAATG 1680
Qy       2044 ACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAGGAGATG 2103
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Db       1681 ACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAGGAGATG 1740
Qy       2104 TTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATCTGTG 2163
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Db       1741 TTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATCTGTG 1800
Qy       2164 TTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACACCGA 2223
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Db       1801 TTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACACCGA 1860
Qy       2224 TCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATTGCAT 2283
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Db       1861 TCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATTGCAT 1920
Qy       2284 CAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGTCTGTTG 2343
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Db       1921 CAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGTCTGTTG 1980
Qy       2344 GCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTATGGCCT 2403
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Db       1981 GCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTATGGCCT 2040
Qy       2404 GGTTTGTCACAGTCCCCATTTCTGGAGTTATCAGTGCTGCCATCATGGCAATCTTCAGAT 2463
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Db       2041 GGTTTGTCACAGTCCCTATTTCTGGAGTTATCAGTGCTGCCATCATGGCAATCTTCAGAT 2100
Qy       2464 ATGTCATCCTCAGAATGTGAAGCTGTTTGAGATTAAAATTTGTGTCAATGTTTGGGACCA 2523
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Db       2101 ATGTCATCCTCAGAATGTGAAGCTGTTTGAGATTAAAATTTGTGTCAATGTTTGGGACCA 2160
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Qy       2524 TCTTAGGTATTCCTGCTCCCCTGAAGAATGATTACAGTGTTAACAGAAGACTGACAAGAG 2583
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Db       2161 TCTTAGGTATTCCTGCTCCCCTGAAGAATGATTACAGTGTTAACAGAAGACTGACAAGAG 2220
Qy       2584 TCTTTTTATTTGGGAGCCAGAGGAGGGAAGTGTTACTTGTGCTATAACTGCTTTTGTGCT 2643
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Db       2221 TCTTTTTATTTGGGAGCCAGAGGAGGGAAGTGTTACTTGTGCTATAACTGCTTTTGTGCT 2280
Qy       2644 AAATATGAATTGTCTCAAAATTAGCTGTGTAAAATAGCCCGGGTTCCACTGGCTCCTGCT 2703
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Qy       2704 GAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCTGTACATATTTCTCTACTTTTTGTATCA 2763
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Qy       2764 GGCTTCAATTCCATTATGTTTTAATGTTGTCTCTGAAGATGACTTGTGATTTTTTTTTCT 2823
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Qy       2824 TTTTTTTAAACCATGAAGAGCCGTTTGACAGAGCATGCTCTGCGTTGTTGGTTTCACCAG 2883
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Qy       2884 CTTCTGCCCTCACATGCACAGGGATTTAACAACAAAAATATAACTACAACTTCCCTTGTA 2943
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Db       2521 CTTCTGCCCTCACATGCACAGGGATTTAACAACAAAAATATAACTACAACTTCCCTTGTA 2580
Qy       2944 GTCTCTTATATAAGTAGAGTCCTTGGTACTCTGCCCTCCTGTCAGTAGTGGCAGGATCTA 3003
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Db       2581 GTCTCTTATATAAGTAGAGTCCTTGGTACTCTGCCCTCCTGTCAGTAGTGGCAGGATCTA 2640
Qy       3004 TTGGCATATTCGGGAGCTTCTTAGAGGGATGAGGTTCTTTGAACACAGTGAAAATTTAAA 3063
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Qy       3064 TTAGTAACTTTTTTGCAAGCAGTTTATTGACTGTTATTGCTAAGAAGAAGTAAGAAAGAA 3123
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Qy       3124 AAAGCCTGTTGGCAATCTTGGTTATTTCTTTAAGATTTCTGGCAGTGTGGGATGGATGAA 3183
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Qy       3184 TGAAGTGGAATGTGAACTTTGGGCAAGTTAAATGGGACAGCCTTCCATGTTCATTTGTCT 3243
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Db       2821 TGAAGTGGAATGTGAACTTTGGGCAAGTTAAATGGGACAGCCTTCCATGTTCATTTGTCT 2880
Qy       3244 ACCTCTTAACTGAATAAAAAAGCCTACAGTTTTTAG 3279
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RESULT 11
US-10-062-674-2102
; Sequence 2102, Application US/10062674
; Publication No. US20040005559A1
; GENERAL INFORMATION:
; APPLICANT: Loring, Jeanne F.; Kaser, Matthew R.
; TITLE OF INVENTION: MARKERS OF NEURONAL DIFFERENTIATION AND MORPHOGENESIS
; FILE REFERENCE: PA-0026-1 CIP
; CURRENT APPLICATION NUMBER: US/10/062,674
; CURRENT FILING DATE: 2002-01-30
; PRIOR APPLICATION NUMBER: US 09/625,102
; PRIOR FILING DATE: 2000-07-24
; NUMBER OF SEQ ID NOS: 2217
; SOFTWARE: PERL Program
; SEQ ID NO 2102
; LENGTH: 3447
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:

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;   NAME/KEY: misc_feature
;   OTHER INFORMATION: Incyte ID No. US20040005559A1 474592.5
;   FEATURE:
;   NAME/KEY: unsure
;   LOCATION: (1) ... (3447)
;   OTHER INFORMATION: a, t, c, g, or other
US-10-062-674-2102

  Query Match             68.3%;  Score 2245.8;  DB 12;  Length 3447;
  Best Local Similarity   91.4%;  Pred. No. 0;
  Matches 3120;  Conservative    0;  Mismatches  170;  Indels  124;  Gaps   65;

Qy          1 CGGCGCTCTCTGCGTGGTTCTTCTTCTCGGCCGCTGAAACCCCCGCGGCTGCTTCCTGGG 60
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Db         34 CGGCGCTCTCTGCGTGGTTCTTCTTCTCGGCCGCTGAAACCCCCGCGGCTGCTTCCTGGG 93

Qy         61 AAGGTCGTGAGTCCCGCTGAGCTGTCCCCGGTGCCGCCGACCCGG-----GCCGTGTGCC 115
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Db         94 AAGGTCGTGAGTCCCGCTGAGCTGTCCCCGGTGCCGCCGACCCGGGCCGTGCCGTGTGCC 153

Qy        116 CGTGGCTCCAGCCGCTGCCGCCTCGATCTCCTCGTCTCCCGCTCCGCCCTCCCTTTTCCC 175
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Db        154 CGTGGCTCCAGCCGCTGCCGCCTCGATCTCCTCGTCTCCCGCTCCGCCCTCCCTTTTCCC 213

Qy        176 T-GGATGAACTTGC-GTCCTTTCTCTTCTCCGCCATGGAATTCT-GCTCCGTGCTTTTAG 232
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Db        214 TGGGATGAACTTGCGGTCCTTTCTCTTCTCCGCCATGGAATTCTGGCTCCGTGGTTTTAG 273

Qy        233 CCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATACGCAGCGTATAGCAGTAAC 292
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        274 CCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATACGCAGCGTATAGCAGTAAC 333

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Db        334 TCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTTATATAATATATATTAT 393

Qy        353 TTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTTGAAAGGCGCTCAGTAG 412
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Db        454 TTCTCTTACTAAACAACCACTACTCCAGAGAATGGCAACGCTGATTACCAGTACTACAGC 513

Qy        473 TGCTACCGCCGCTTCTGGTCCTTTGGTGGACTACCTATGGATGCTCATCCTGGGCTTCAT 532
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Qy        533 TATTGCATTTGTCTTG-GCATTCTCCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGT- 590
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Db        574 TATTGCATTTGTCTTGTGCATTCTCCGTGGGAGCCAATGATGTAGCAAAGTTCTTTTGTG 633

Qy        591 ------ACAGCTGTGGGCTCAGGTGTAGTGACCCTGAA---GCAAGCCTGCATCCTAGCT 641
                  || | |||||||||| ||||||||||||||||     ||||||||||||||||||
Db        634 TACACGACTGGTGTGGGCTCANGTGTAGTGACCCTGAAAGCAAAAGCCTGCATCCTAGCT 693

Qy        642 AGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCATCCGG 701
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Db        694 AGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAACCATCCGG 753

Qy        702 AAGGGCTTGAT-TGACGTGGAGATGTACAACTC---GACTCAAGGGCTACTG-ATGGCCG 756
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Db        754 AAGGGCTTGATATGACGTGGAGATGTACAACTCCGACTCCAAGGGGCTACTGAATGGCCG 813

Qy        757 GCTCAGTCA--GTGCTATGTTTGGTTCT--GCTGTGTGGCAACTCGTGGCTTCGTTTTTG 812
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Db        814 GCTCAGTCATGTTGCTATGTTTGGTTCTTGCTGTGGTGGCAACTCGTGGCTTCGTTTTTG 873

Qy        813 AAGCTCCC---TATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGG---TTTCTCC 866
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Db        874 AAGCTCCCCTATTTCTGGGAACCCATTTGTATTGTTGGTGCAACTATTGGGTTTCTCCCC 933

Qy        867 CTCGTGGCAAAGGGGCAGGAGGGTGTC-AAGTGGTCTGAACTGATAAAAATTGTGATGTC 925
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Db        934 TCGTGGCAAAAGGGGCAGGAGGGTGTCAAAGTGGTCTGAACTGATAAAAATTGTGATGTC 993

Qy        926 TTGGTT-CGTGTCCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGTTC 984
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Qy        985 GTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTT 1044
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Db       1054 GTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTT 1113

Qy       1045 TCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAGCACCGTTGC 1104
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Qy       1105 TGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGTTT 1164
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Db       1174 TGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGTTT 1233

Qy       1165 TCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAATTGAACGAG 1224
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Db       1234 TCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAATTGAACGAG 1293

Qy       1225 AAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAG 1284
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1294 AAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAG 1353

Qy       1285 ACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGAGG 1344
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Db       1354 ACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGAGG 1413

Qy       1345 --TAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCTCATT-CA 1401
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Db       1414 GTTAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGGCTCCTTCCA 1473

Qy       1402 AACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGA-GGCTTCCCAGCGTGGACTTGAA- 1459
              |||||||||||||||||||||||||||||||||||| |||||||||||||||||||||| 
Db       1474 AACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGGCTTCCCAGCGTGGACTTGAAC 1533

Qy       1460 AGAGGAAACCAGC---ATAGATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGG-- 1514
              |||||||||||||     |||||||||||||||||||||||||||||||||||||||| 
Db       1534 AGAGGAAACCAGNTCACTAGATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGGA 1593

Qy       1515 AACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTGGCCACTACCAG 1574
              | | |  ||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1594 ACCTTGTCCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTGGCCACTACCAG 1653

Qy       1575 TATCACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTT 1634
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1654 TATCACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTT 1713

Qy       1635 GCCAAGGT-GGGAGATTGCATGGGAGACTCCGGTGACAAACCCTTAAGGCGCAATAATAG 1693
              |||||||| |||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1714 GCCAAGGTGGGGAGATTGCATGGGAGACTCCGGTGACAAACCCTTAAGGCGCAATAATAG 1773

Qy       1694 CTATACTTCCTATACCATGGCAATATGT-GGCATGCCTCTGGATTCATTCCGTGCCAAAG 1752
              |||||||||||||||||||||||||||| |||||||||||||||||||||||||||||||
Db       1774 CTATACTTCCTATACCATGGCAATATGTGGGCATGCCTCTGGATTCATTCCGTGCCAAAG 1833

Qy       1753 AAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATGCAGACTCCAAGA 1812
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1834 AAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATGCAGACTCCAAGA 1893

Qy       1813 AGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACCTTCACTCAG 1872
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1894 AGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACCTTCACTCAG 1953

Qy       1873 CATCTGAGATAGACATGAGTGTCAAGGCAGAG-ATGGGTCTAGGTGACAGAAAAGGAAGT 1931
              |||||||||||||||||||||||||||||||| |||||||||||||||||||||||||||
Db       1954 CATCTGAGATAGACATGAGTGTCAAGGCAGAGTATGGGTCTAGGTGACAGAAAAGGAAGT 2013

Qy       1932 AATGGCTCTCTAGAAGAATGG-TATGACCAGGATAAGCCTGAAGTCTCTCT---CCTCTT 1987
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Db       2014 AATGGCTCTCTAGAAGAATGGTTATGACCAGGATAAGCCTGAAGTACTACTCTACCTCTT 2073

Qy       1988 CCAGTTCCTG-CAGATCCTTACAGCCTGCTTTGGGTCATTCG---CCCATGGTGGCAATG 2043
              |||||||||| |||||||||||||||||||| ||    | |   ||||||||||||||||
Db       2074 CCAGTTCCTGCCAGATCCTTACAGCCTGCTTTTGGGTCATTGCGACCCATGGTGGCAATG 2133

Qy       2044 -ACGTAAGCAATGCCATTGGGCCTCTGGTT-GCTTTATATTTGGTTTATGACACAGGAGA 2101
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Db       2134 CACGTAAGCAATGCCATTGGGCCTCTGGTTGGCTTTATATTTGGTTTATGACACAGGAGA 2193

Qy       2102 TGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATCTG 2161
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Db       2194 TGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATCTG 2253

Qy       2162 TGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAG--ACCATGGGGAAGGATCTGACA 2219
              |||||||||||||||||||||||||||| |||||||   |   ||| ||||||||||||
Db       2254 TGTTGGTCTGTGGGTTTGGGGAAGAAGAGGTATCCAGGACCATGGGGCAAGGATCTGACA 2313

Qy       2220 CCGATCACACCCTCTAGTGGCTT---CAGTATTGAACTGGCATCTGCCC-TCACTGTGGT 2275
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Db       2314 CCGATCACACCCTCTAGCTGGCTTCCAGTATTTGAACTGCAATCTGCCCTTCACTGTGGT 2373

Qy       2276 GATTGCATC-AAATATTGGCCTTCCCATCAG---TACAACACATTGTAAA----GTGGGC 2327
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Db       2374 GATTGCATCAAAATATTGGCCTTCCCATCAGCTTACAATCACATTGTAAAGGCTGGGCCC 2433

Qy       2328 TCTGTTGTGT-CTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCG 2386
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Db       2434 TCTGTTGTGTCCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCN 2493

Qy       2387 TAACATTTTT-----ATGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCAGTGCT 2441
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Db       2494 GTTAACATTTTCTACTGGCCTGGTTTCGTCACAGTCCCCATTTCTGGAGTTATCAGTGCT 2553

Qy       2442 GCCAT-CATGGCAATCTTCAGATATGTCAT-CCTCAGAATGTGAAGCTGTTTGAGATTAA 2499
              ||||| |||||||||||||||||||||||| |||||||||||||||||||||||||||||
Db       2554 GCCATACATGGCAATCTTCAGATATGTCATCCCTCAGAATGTGAAGCTGTTTGAGATTAA 2613

Qy       2500 AATTTGTGTCAATGTTTGGGACCATCTT-AGGTATTCCTGCTCCCCTGAAGAATGA-TTA 2557
              |||||||||||||||||||||||||||| |||||||||||||||||||||||||| |||
Db       2614 AATTTGTGTCAATGTTTGGGACCATCTTAAGGTATTCCTGCTCCCCTGAAGAATGATTTA 2673

Qy       2558 CAGTGTTAACAG-AAGACTGACA-AGAGTCTTTTTATTTGGGAGCCAGAGGAGGGAAGTG 2615
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Db       2854 AATTCCTGTACATATTTCTCTACTTTTTGTATCAGGGCTTCAATTCCATTATGTTCTTAA 2913

Qy       2788 TGTTGTCTCTG--AAGATGACTTGTGATTTTTTTTT--CTTTTTTTTAAACCATGAAGAG 2843
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Db       2914 TGTTGTCTCTGGAAGGATGACTTGTGATTTTTTTTTTCTTTTTTTTAAACCATGGAAGAG 2973

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              |||||||||||| |||||| ||| ||||||||||||||    | |     ||  |||
Db       2974 CCGTTTGACAGAGGCATGCGCTGCCGTTGTTGGTTTCACCCAGCTTTCTGGCCCTCAACA 3033

Qy       2898 TGCACAGGGATTTAAC--AACAAAAATATAACTACAACTT--CCCTTGTAGTCTCTTATA 2953
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Db       3034 TGCACAGGGATTTAACCAACAAAAATTATAACTACAACTTACCCCTTGTAGTCCCCCTTA 3093

Qy       2954 T---AAGTAGAGTCCTTGGTACTCTG---CCCTCCTGTCAGTAGTGGCAGG---ATCTAT 3004
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Db       3094 TATAAGTTAGAGTCCCTTGGTACCTCTGCCCCTCCTGTCAGTAGTGGGCAGGATCCTATT 3153

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Db       3154 GGGCATATTCGGGAGCCTTCCTTAGAGGGGATGAGGTTCCTTTGAACACAGGTGAAAATT 3213

Qy       3060 TAAATTAGTAAC-TTTTTTGCAAGCAGTTTATTGACTGTTATTGCTAAGAAG-AAGTAAG 3117
              |||||||||||| |||||||||||||||||||||||||||||||||||||| |||||||
Db       3214 TAAATTAGTAACTTTTTTTGCAAGCAGTTTATTGACTGTTATTGCTAAGAAGAAAGTAAG 3273

Qy       3118 AAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAAGATTTCTGGCAGTGTGGGATG 3177
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Db       3274 AAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAAGATTTCTGGCAGTGTGGGATG 3333

Qy       3178 GATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAATGGGACAGCCTTCCATGTTCAT 3237
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3334 GATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAATGGGACAGCCTTCCATGTTCAT 3393

Qy       3238 TTGTCTACCTCTTAACTGAATA-AAAAAGCCTACAGTTTTTAGAAAAAAAAAAA 3290
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RESULT 12
US-10-314-790-4
; Sequence 4, Application US/10314790
; Publication No. US20040002079A1
;   GENERAL INFORMATION:
;        APPLICANT: EMORY UNIVERSITY
;                   Gunn, Robert B.
;                   Timmer, Richard T.
;        TITLE OF INVENTION: SODIUM-PHOSPHATE COTRANSPORTER IN
;                            LITHIUM THERAPY FOR THE TREATMENT OF MENTAL ILLNESS
;        NUMBER OF SEQUENCES: 12
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: Kilpatrick Stockton LLP
;             STREET: Suite 2800, 1100 Peachtree Street NE
;             CITY: Atlanta
;             STATE: GA
;             COUNTRY: USA
;             ZIP: 30309-4530
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: Floppy disk
;             COMPUTER: IBM PC compatible
;             OPERATING SYSTEM: PC-DOS/MS-DOS
;             SOFTWARE: PatentIn Release #1.0, Version #1.25
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/10/314,790
;             FILING DATE: 09-Dec-2002
;             CLASSIFICATION: <Unknown>
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: US/09/380,164
;             FILING DATE: <Unknown>
;             APPLICATION NUMBER: PCT/US98/02875
;             FILING DATE: <Unknown>
;        ATTORNEY/AGENT INFORMATION:
;             NAME: Meredith, Roy D.
;             REGISTRATION NUMBER: 30,777
;             REFERENCE/DOCKET NUMBER: EMU153PCT
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: (404) 815-6500
;             TELEFAX: (404) 815-6555
;   INFORMATION FOR SEQ ID NO: 4:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 3175 base pairs
;             TYPE: nucleic acid
;             STRANDEDNESS: single
;             TOPOLOGY: linear
;        MOLECULE TYPE: mRNA
;        HYPOTHETICAL: NO
;        ANTI-SENSE: NO
;        ORIGINAL SOURCE:
;             TISSUE TYPE: placenta
;        FEATURE:
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;             NAME/KEY:  hPiT-2
;             LOCATION:  1..3175
;             OTHER INFORMATION:  /product= "Leukemia virus receptor 2"
;             /label= Receptor2
;         PUBLICATION INFORMATION:
;             DOCUMENT NUMBER:  van Zeijl, M., et al., "A Human Amphotropic Retrovirus R
;         PUBLICATION INFORMATION:
;             DOCUMENT NUMBER:  US 5,550,221 P
;         SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-10-314-790-4

  Query Match            18.8%;  Score 617.2;  DB 12;  Length 3175;
  Best Local Similarity  60.3%;  Pred. No. 2.8e-167;
  Matches 1203;  Conservative    0;  Mismatches  718;  Indels   75;  Gaps    8;

Qy        496 TGGTGGACTACCTATGGATGCTCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCT 555
Db        251 TGGATGAGTATTTGTGGATGGTCATTTTGGGTTTCATCATAGCTTTCATCTTGGCCTTTT 310

Qy        556 CCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGA 615
Db        311 CTGTTGGTGCAAACGATGTTGCCAACTCCTTTGGTACAGCCGTGGGCTCTGGTGTGGTGA 370

Qy        616 CCCTGAAGCAAGCCTGCATCCTAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGG 675
Db        371 CCTTGAGGCAGGCATGCATTTTAGCTTCAATATTTGAAACCACCGGCTCCGTGTTACTAG 430

Qy        676 GGGCCAAAGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGA 735
Db        431 GCGCCAAAGTAGGAGAAACCATTCGCAAAGGTATCATTGACGTGAACCTGTACAACGAGA 490

Qy        736 CTCAAGGGCTACTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAAC 795
Db        491 CGGTGGAGACTCTCATGGCTGGGGAAGTTAGTGCCATGGTTGGTTCCGCTGTGTGGCAGC 550

Qy        796 TCGTGGCTTCGTTTTTGAAGCTCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTA 855
Db        551 TGATTGCTTCCTTCCTGAGGCTTCCAATCTCAGGAACGCACTGCATTGTGGGTTCTACTA 610

Qy        856 TTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAA 915
Db        611 TAGGATTCTCACTGGTCGCAATCGGTACCAAAGGTGTGCAGTGGATGGAGCTTGTCAAGA 670

Qy        916 TTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCT 975
Db        671 TTGTTGCTTCTTGGTTTATATCTCCACTGTTGTCTGGTTTCATGTCTGGCCTGCTGTTTG 730

Qy        976 TCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTT 1035
Db        731 TACTCATCAGAATTTTCATCTTAAAAAAGGAAGACCCTGTTCCCAATGGCCTCCGGGCAC 790

Qy       1036 TGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAG 1095
Db        791 TCCCAGTATTCTATGCTGCTACCATAGCAATCAATGTCTTTTCCATCATGTACACAGGAG 850

Qy       1096 CACCGTTGCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGAT 1155
Db        851 CACCAGTGCTCGGCCTTG---TTCTCCCCATGTGGGCCATAGCCCTCATTTCCTTTGGTG 907

Qy       1156 GTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAA 1215
Db        908 TCGCCCTCCTGTTCGCTTTTTTTGTGTGGCTCTTCGTGTGTCCGTGGATGCGGAGGAAAA 967

Qy       1216 TTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCT 1275
Db        968 TAACAGGCAAATTACAAAAAGAAGGTGCTTTATCACGAGTATCTGACGAAAGCCTCAGTA 1027

Qy       1276 TGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTG 1335
Db       1028 AGGTTCAGGAAGCAGAGTCCCCAGTATT-----------------TAAAGAGCTACCAGG 1070

Qy       1336 TTTCTGAGGTAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCT 1395
Db       1071 TGCCAAGGCTAATGATGACAGCACCATCCCGCTCACGGGAGCAGCAGGGGAG-------- 1122

Qy       1396 CATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACT 1455
Db       1123 -----ACACTGGGGACCTCGGAAGGCACTTCTGCG-------GGCAGCCACCCTCGGGCT 1170

Qy       1456 TGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGA 1515
Db       1171 GCATACGGAAGAGCACTGTCCATGACCCATGGCTCTGTGAAATCGCCCATCTCCAACGGC 1230

Qy       1516 ACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTGGCCACTACCAGT 1575
Db       1231 ACCTT-----------------CGGCTTCGACGGCCACACCAGGAGCGACGGTCATGTGT 1273

Qy       1576 ATCACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTTG 1635
Db       1274 ACCACACCGTGCACAAAGACTCGGGGCTCTACAAAGATCTGCTGCACAAAATCCACATCG 1333

Qy       1636 CCAAGGTGGGAGATTGCATGGGAGACTCCGGTGACAA------ACCCTTAAGGCGCAATA 1689
Db       1334 ACAGGGGCCCCGAGGAGAAGCCAGCCCAGGAAAGCAACTACCGGCTGCTCCGCCGAAACA 1393

Qy       1690 ATAGCTATACTTCCTATACCATGGCAATATGTGGCATGCCTCTGGATTCATTCCGTGCCA 1749
Db       1394 ACAGTTACACCTGCTACACCGCAGCCATTTGTGGGCTGCCAGTGCACGCCACCTTTCGAG 1453

Qy       1750 AAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATG---CAGACT 1806
Db       1454 CTGCGGACTCATCGGCCCCAGAGGACAGTGAGAAGCTGGTGGGCGACACCGTGTCCTACT 1513

Qy       1807 CCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACCTTC 1866
Db       1514 CCAAGAAGAGGCTGCGCTACGACAGCTACTCGAGCTACTGTAACGCGGTGGCAGAGGCGG 1573

Qy       1867 ACTCAGCATCTGAGA---------TAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTG 1917
Db       1574 AGATCGAGGCGGAGGAGGGCGGCGTGGAGATGAAGCTGGCGTCGGAGCTGGCCGACCCTG 1633

Qy       1918 ACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATGACCAGGATAAGCCTGAAGTCT 1977
Db       1634 ACCAGCCGCGAGAGGACCCTGCAGAGGAGGAGAAGGAGGAGAAGGACGCACCCGAGGTTC 1693

Qy       1978 CTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTG 2037
Db       1694 ACCTCCTGTTCCATTTCCTGCAGGTCCTCACCGCCTGTTTCGGGTCCTTTGCTCACGGCG 1753

Qy       2038 GCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAG 2097
Db       1754 GCAATGACGTGAGTAATGCCATCGGTCCCCTGGTAGCCTTGTGGCTGATTTACAAACAAG 1813

Qy       2098 GAGATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTA 2157
Db       1814 GCGGGGTAACGCAAGAAGCAGCTACACCCGTCTGGCTGCTGTTTTATGGAGGAGTTGGAA 1873

Qy       2158 TCTGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGA 2217
Db       1874 TCTGCACAGGCCTCTGGGTCTGGGGGAGAAGAGTGATCCAGACCATGGGGAAGGACCTCA 1933

Qy       2218 CACCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGA 2277
Db       1934 CTCCCATCACGCCGTCCAGCGGCTTCACGATCGAGCTGGCCTCAGCCTTCACAGTGGTGA 1993

Qy       2278 TTGCATCAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGT 2337
Db       1994 TCGCCTCCAACATCGGGCTTCCAGTCAGCACCACGCACTGTAAGGTGGGCTCGGTGGTGG 2053

Qy       2338 CTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTA 2397
Db       2054 CCGTGGGCTGGATCCGCTCCCGCAAGGCTGTGGACTGGCGCCTCTTTCGGAACATCTTCG 2113

Qy       2398 TGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCAGTGCTGCCATCATGGCAATCT 2457
Db       2114 TGGCCTGGTTCGTGACCGTCCCTGTGGCTGGGCTGTTCAGCGCTGCTGTCATGGCTCTTC 2173
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Qy       2458 TCAGATATGTCATCCT 2473
Db       2174 TCATGTATGGGATCCT 2189

RESULT 13
US-09-873-367C-143
; Sequence 143, Application US/09873367C
; Publication No. US20030165839A1
; GENERAL INFORMATION:
;  APPLICANT: Young, Paul
;  APPLICANT:  Soppet, Daniel
;  APPLICANT:  Endress, Gregory
;  APPLICANT:  Augustus, Meena
;  APPLICANT:  Ebner, Reinhard
;  APPLICANT:  Carter, Kenneth
;  TITLE OF INVENTION: Cancer Gene Determination and Therapeutic Screening Using
;  TITLE OF INVENTION:  Signature Gene Sets
;  FILE REFERENCE: 689290-64
;  CURRENT APPLICATION NUMBER: US/09/873,367C
;  CURRENT FILING DATE:  2003-04-29
;  PRIOR APPLICATION NUMBER: U.S. 60/236,891
;  PRIOR FILING DATE: 2000-09-29
;  PRIOR APPLICATION NUMBER: U.S. 60/236,842
;  PRIOR FILING DATE: 2000-09-29
;  PRIOR APPLICATION NUMBER: U.S. 60/244,867
;  PRIOR FILING DATE: 2000-11-01
;  PRIOR APPLICATION NUMBER: U.S. 60/245,084
;  PRIOR FILING DATE: 2000-11-01
;  NUMBER OF SEQ ID NOS: 1067
;  SOFTWARE: PatentIn version 3.0
; SEQ ID NO 143
;   LENGTH: 3175
;   TYPE: DNA
;   ORGANISM: Homo sapiens
US-09-873-367C-143

  Query Match             18.8%;  Score 617.2;  DB 13;  Length 3175;
  Best Local Similarity   60.3%;  Pred. No. 2.8e-167;
  Matches 1203;  Conservative    0;  Mismatches  718;  Indels   75;  Gaps    8;

Qy        496 TGGTGGACTACCTATGGATGCTCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCT 555
Db        251 TGGATGAGTATTTGTGGATGGTCATTTTGGGTTTCATCATAGCTTTCATCTTGGCCTTTT 310

Qy        556 CCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGA 615
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Qy        856 TTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAA 915
Db        611 TAGGATTCTCACTGGTCGCAATCGGTACCAAAGGTGTGCAGTGGATGGAGCTTGTCAAGA 670

Qy        916 TTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCT 975
Db        671 TTGTTGCTTCTTGGTTTATATCTCCACTGTTGTCTGGTTTCATGTCTGGCCTGCTGTTTG 730

Qy        976 TCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTT 1035
Db        731 TACTCATCAGAATTTTCATCTTAAAAAAGGAAGACCCTGTTCCCAATGGCCTCCGGGCAC 790

Qy       1036 TGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAG 1095
Db        791 TCCCAGTATTCTATGCTGCTACCATAGCAATCAATGTCTTTTCCATCATGTACACAGGAG 850

Qy       1096 CACCGTTGCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGAT 1155
Db        851 CACCAGTGCTCGGCCTTG---TTCTCCCCATGTGGGCCATAGCCCTCATTTCCTTTGGTG 907

Qy       1156 GTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAA 1215
Db        908 TCGCCCTCCTGTTCGCTTTTTTTGTGTGGCTCTTCGTGTGTCCGTGGATGCGGAGGAAAA 967

Qy       1216 TTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCT 1275
Db        968 TAACAGGCAAATTACAAAAAGAAGGTGCTTTATCACGAGTATCTGACGAAAGCCTCAGTA 1027

Qy       1276 TGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTG 1335
Db       1028 AGGTTCAGGAAGCAGAGTCCCCAGTATT-----------------TAAAGAGCTACCAGG 1070

Qy       1336 TTTCTGAGGTAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCT 1395
Db       1071 TGCCAAGGCTAATGATGACAGCACCATCCCGCTCACGGGAGCAGCAGGGGAG-------- 1122

Qy       1396 CATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACT 1455
Db       1123 -----ACACTGGGGACCTCGGAAGGCACTTCTGCG-------GGCAGCCACCCTCGGGCT 1170

Qy       1456 TGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGA 1515
Db       1171 GCATACGGAAGAGCACTGTCCATGACCCATGGCTCTGTGAAATCGCCCATCTCCAACGGC 1230

Qy       1516 ACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTGGCCACTACCAGT 1575
Db       1231 ACCTT-----------------CGGCTTCGACGGCCACACCAGGAGCGACGGTCATGTGT 1273

Qy       1576 ATCACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTTG 1635.
Db       1274 ACCACACCGTGCACAAAGACTCGGGGCTCTACAAAGATCTGCTGCACAAAATCCACATCG 1333

Qy       1636 CCAAGGTGGGAGATTGCATGGGAGACTCCGGTGACAA------ACCCTTAAGGCGCAATA 1689
Db       1334 ACAGGGGCCCCGAGGAGAAGCCAGCCCAGGAAAGCAACTACCGGCTGCTCCGCCGAAACA 1393

Qy       1690 ATAGCTATACTTCCTATACCATGGCAATATGTGGCATGCCTCTGGATTCATTCCGTGCCA 1749
Db       1394 ACAGTTACACCTGCTACACCGCAGCCATTTGTGGGCTGCCAGTGCACGCCACCTTTCGAG 1453

Qy       1750 AAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATG---CAGACT 1806
Db       1454 CTGCGGACTCATCGGCCCCAGAGGACAGTGAGAAGCTGGTGGGCGACACCGTGTCCTACT 1513

Qy       1807 CCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACCTTC 1866
Db       1514 CCAAGAAGAGGCTGCGCTACGACAGCTACTCGAGCTACTGTAACGCGGTGGCAGAGGCGG 1573

Qy       1867 ACTCAGCATCTGAGA---------TAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTG 1917
Db       1574 AGATCGAGGCGGAGGAGGGCGGCGTGGAGATGAAGCTGGCGTCGGAGCTGGCCGACCCTG 1633

Qy       1918 ACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATGACCAGGATAAGCCTGAAGTCT 1977
Db       1634 ACCAGCCGCGAGAGGACCCTGCAGAGGAGGAGAAGGAGGAGAAGGACGCACCCGAGGTTC 1693

Qy       1978 CTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTG 2037
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Qy       2038 GCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAG 2097
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Qy       2098 GAGATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTA 2157
Db       1814 GCGGGGTAACGCAAGAAGCAGCTACACCCGTCTGGCTGCTGTTTTATGGAGGAGTTGGAA 1873

Qy       2158 TCTGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGA 2217
Db       1874 TCTGCACAGGCCTCTGGGTCTGGGGGAGAAGAGTGATCCAGACCATGGGGAAGGACCTCA 1933

Qy       2218 CACCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGA 2277
Db       1934 CTCCCATCACGCCGTCCAGCGGCTTCACGATCGAGCTGGCCTCAGCCTTCACAGTGGTGA 1993

Qy       2278 TTGCATCAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGT 2337
Db       1994 TCGCCTCCAACATCGGGCTTCCAGTCAGCACCACGCACTGTAAGGTGGGCTCGGTGGTGG 2053

Qy       2338 CTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTA 2397
Db       2054 CCGTGGGCTGGATCCGCTCCCGCAAGGCTGTGGACTGGCGCCTCTTTCGGAACATCTTCG 2113

Qy       2398 TGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCAGTGCTGCCATCATGGCAATCT 2457
Db       2114 TGGCCTGGTTCGTGACCGTCCCTGTGGCTGGGCTGTTCAGCGCTGCTGTCATGGCTCTTC 2173

Qy       2458 TCAGATATGTCATCCT 2473
Db       2174 TCATGTATGGGATCCT 2189


RESULT 14
US-10-388-934-200
; Sequence 200, Application US/10388934
; Publication No. US20040005547A1
; GENERAL INFORMATION:
;  APPLICANT: Boess, Franziska
;  APPLICANT:  Suter-Dick, Laura
;  APPLICANT:  Wolf, Detlef
;  TITLE OF INVENTION: BIOMARKERS AND EXPRESSION PROFILES FOR TOXICOLOGY
;  FILE REFERENCE: 21199
;  CURRENT APPLICATION NUMBER: US/10/388,934
;  CURRENT FILING DATE:  2003-03-14
;  PRIOR APPLICATION NUMBER: 02005336.9
;  PRIOR FILING DATE: 2002-03-14
;  PRIOR APPLICATION NUMBER: 02015657.6
;  PRIOR FILING DATE: 2002-07-17
;  NUMBER OF SEQ ID NOS: 862
;  SOFTWARE: PatentIn version 3.1
; SEQ ID NO 200
;   LENGTH: 2287
;   TYPE: DNA
;   ORGANISM: Rattus norvegicus (No. US20040005547A1way rat)
US-10-388-934-200

  Query Match             18.7%;  Score 615;  DB 12;  Length 2287;
  Best Local Similarity   60.0%;  Pred. No. 9.5e-167;
  Matches 1195;  Conservative    0;  Mismatches  720;  Indels   78;  Gaps    7;

Qy        496 TGGTGGACTACCTATGGATGCTCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCT 555
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Qy        556 CCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGA 615
Db        338 CAGTTGGTGCCAATGATGTTGCCAACTCCTTTGGAACCGCGGTGGGCTCTGGCGTGGTGA 397

Qy        616 CCCTGAAGCAAGCCTGCATCCTAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGG 675
Db        398 CCCTGAGACAGGCATGCATTCTGGCTTCGATATTTGAAACCACCGGCTCTGTGCTTTTGG 457

Qy        676 GGGCCAAAGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGA 735
Db        458 GAGCGAAAGTGGGAGAGACTATCCGGAAGGGCATCATTGACGTGAACCTCTACAATGAGA 517

Qy        736 CTCAAGGGCTACTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAAC 795
Db        518 CTGTGGAAACGCTCATGGCTGGGGAAGTGAGTGCAATGGTCGGCTCAGCGGTCTGGCAGT 577

Qy        796 TCGTGGCTTCGTTTTTGAAGCTCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTA 855
Db        578 TGATCGCGTCCTTCCTGAGACTCCCAATCTCAGGGACACACTGCATCGTGGGCTCTACCA 637

Qy        856 TTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAA 915
Db        638 TTGGCTTCTCGCTCGTGGCCATCGGCCCAAAGGGAGTGCAGTGGATGGAACTCGTCAAGA 697

Qy        916 TTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCT 975
Db        698 TCGTTGCCTCCTGGTTCATATCCCCACTTCTGTCTGGCTTCATGTCTGGTGTGCTGTTCA 757

Qy        976 TCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTT 1035
Db        758 TCCTGATCAGAATGTTCATCCTAACCAAGGAGGACCCTGTTCCAAACGGTCTCCAGGCAC 817

Qy       1036 TGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAG 1095
Db        818 TTCCTCTGTTTTATGCTGCCACTATAGCAATAAACGTCTTCTCCATCATGTACACAGGAG 877

Qy       1096 CACCGTTGCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGAT 1155
Db        878 CACCAGTGCTGGGCCT---GTCGCTCCCCATCTGGGCAATAGCGCTCATCTCCTTCGGCG 934

Qy       1156 GTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAA 1215
Db        935 TTGCCCTGCTGTTCGCCTTCTTCGTGTGGCTGTTCGTGTGTCCGTGGATGAAGAGGAAAA 994

Qy       1216 TTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCT 1275
Db        995 TAGCGGGCAGGTTAGAGAAGGAGAGTGCTCTGTCCCGCGCGTCTGACGAAAGCCTCAGGA 1054

Qy       1276 TGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTG 1335
Db       1055 AGGTACAGGAAGCTGAGTCCCCTGTCTTTAAGGAGCTTCCTGGTGCCAAAGCCAGTGATG 1114

Qy       1336 TTTCTGAGGTAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCT 1395
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Qy       1396 CATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACT 1455
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Qy       1456 TGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGA 1515
Db       1198 TCCTACGGCCGAGCACTTTCCATGACTCACGGCTCTGCAAAGTCACCCATCTCCAATGGA 1257

Qy       1516 ACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTGGCCACTACCAGT 1575
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Db       1301 ATCATACTGTACACAAAGACTCGGGACTCTACAAAGACTTGCTGCACAAAATCCATGTAG 1360

Qy       1636 CCAAGGTGGGAGATTGCATGGGA------GACTCCGGTGACAAACCCTTAAGGCGCAATA 1689
Db       1361 ACAAGGGCCCCGAGGAGAAACCAGCTCAGGAAAACAACTACAGGCTGCTCCGGAGGAACA 1420

Qy       1690 ATAGCTATACTTCCTATACCATGGCAATATGTGGCATGCCTCTG---GATTCATTCCGTG 1746
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Qy       1915 GTGACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATGACCAGGATAAGCCTGAAG 1974

Db       1661 CTGACCAGCCTCACGAGGATCCTGCGGAAGACGAGAAGGAGGAGAAGGACTCCGCAGAGG 1720

Qy       1975 TCTCTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATG 2034

Db       1721 TCCACCTGCTCTTCCACTTCCTGCAAGTCCTCACTGCCTGCTTTGGGTCCTTTGCTCACG 1780

Qy       2035 GTGGCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACA 2094

Db       1781 GTGGCAACGATGTGAGCAATGCTATCGGTCCCCTTGTGGCCCTGTGGCTGATATATAAAC 1840

Qy       2095 CAGGAGATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTG 2154

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Qy       2275 TGATTGCATCAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTG 2334

Db       2021 TGATCGCCTCCAACATTGGGCTTCCCGTCAGCACCACACACTGCAAGGTGGGCTCAGTGG 2080

Qy       2335 TGTCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTT 2394

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Qy       2455 TCTTCAGATATGT 2467

Db       2201 TCCTCATGTATAT 2213


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; Sequence 623, Application US/09879536
; Patent No. US20020144298A1
; GENERAL INFORMATION:
;  APPLICANT: Endege, Wilson O.
;  APPLICANT:  Steinmann, Kathleen E.
;  APPLICANT:  Astle, Jon H.
;  APPLICANT:  Burgess, Christopher C.
;  APPLICANT:  Bushnell, Steven E.
;  APPLICANT:  Carroll III, Eddie
;  APPLICANT:  Catino, Theodore J.
;  APPLICANT:  Derti, Adnan
;  APPLICANT:  Ford, Donna M.
;  APPLICANT:  Lewis, Marcia E.
;  APPLICANT:  Monahan, John E.
;  APPLICANT:  Schlegel, Robert
;  TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
;  TITLE OF INVENTION:  PRODUCTS
;  FILE REFERENCE: CCD-257 (US)
;  CURRENT APPLICATION NUMBER: US/09/879,536
;  CURRENT FILING DATE:  2001-09-21
;  PRIOR APPLICATION NUMBER: US 60/088,801
;  PRIOR FILING DATE: 1998-06-10
;  NUMBER OF SEQ ID NOS: 850
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 623
;   LENGTH: 662
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(662)
;   OTHER INFORMATION: n = A,T,C or G
US-09-879-536-623

  Query Match            15.1%;  Score 495.2;  DB 10;  Length 662;
  Best Local Similarity  90.2%;  Pred. No. 2.1e-132;
  Matches  591;  Conservative    0;  Mismatches   54;  Indels   10;  Gaps    6;

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Qy       1726 TGCCTCTGGATTCATTCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGC 1785

Db        121 TGCCTCTGGATTCATTCCGTGCCAAAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGC 180

Qy       1786 TGACATGGCCTAATGCAGACTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACT 1845

Db        181 TGACATGGCCTAATGCGGACTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACT 240

Qy       1846 GCAATGCTGTGTCTGACCTTCACTCAGCATCTGAGATAGACATGAGTGTCAAGGCAGAGA 1905

Db        241 GCAATGCTGTGTCTGACCTTCACTCAGCATCTGAGATAGACATGAGTGTCAAGGCAGAGA 300

Qy       1906 TGGGTCTAGGTGACAGAAAAGGAA--GTAATGGCTCTCTAGAAGAATGGTATGACCAGGA 1963

Db        301 TGGGTCTAGGTGACAGAAAAGGAAAGTAATGGGCTCTCTAGAAGAATGGTATGACCAGGA 360

Qy       1964 TAAGCCTGAAGTCTCTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTC 2023

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Qy       2142 TAT--GGTGGTGTTGGTATCTGTGTTGG---TCTGTGGGTTTGGGGAAGAAGAGTTATCC 2196

Db        540 TANTGGNGGGGGTTGGGATCTGNGGTTGGTCTGTGGGGTTTGGGGAAAAAAAAGTTTTCC 599

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GenCore version 5.1.6

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Perfect score: 3290
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Gapop 10.0 , Gapext 1.0

Searched: 569978 seqs, 220691566 residues

Total number of hits satisfying chosen parameters: 1139956

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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	2	3172	96.4	3211	2	US-08-436-900A-1	Sequence 1, Appli
	3	2133	64.8	3260	1	US-07-674-287B-3	Sequence 3, Appli
	4	2131.4	64.8	3260	2	US-08-436-900A-3	Sequence 3, Appli
	5	617.2	18.8	3175	1	US-08-050-684-1	Sequence 1, Appli
	6	617.2	18.8	3175	1	US-08-582-719-1	Sequence 1, Appli
	7	495.2	15.1	662	3	US-09-328-111-623	Sequence 623, App
	8	451.4	13.7	613	3	US-09-328-111-521	Sequence 521, App
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	11	99.8	3.0	1830121	4	US-09-643-990A-1	Sequence 1, Appli
	12	87	2.6	1380	4	US-09-252-991A-6471	Sequence 6471, Ap
	13	87	2.6	1407	4	US-09-252-991A-6303	Sequence 6303, Ap
c	14	86	2.6	1230025	4	US-09-198-452A-1	Sequence 1, Appli
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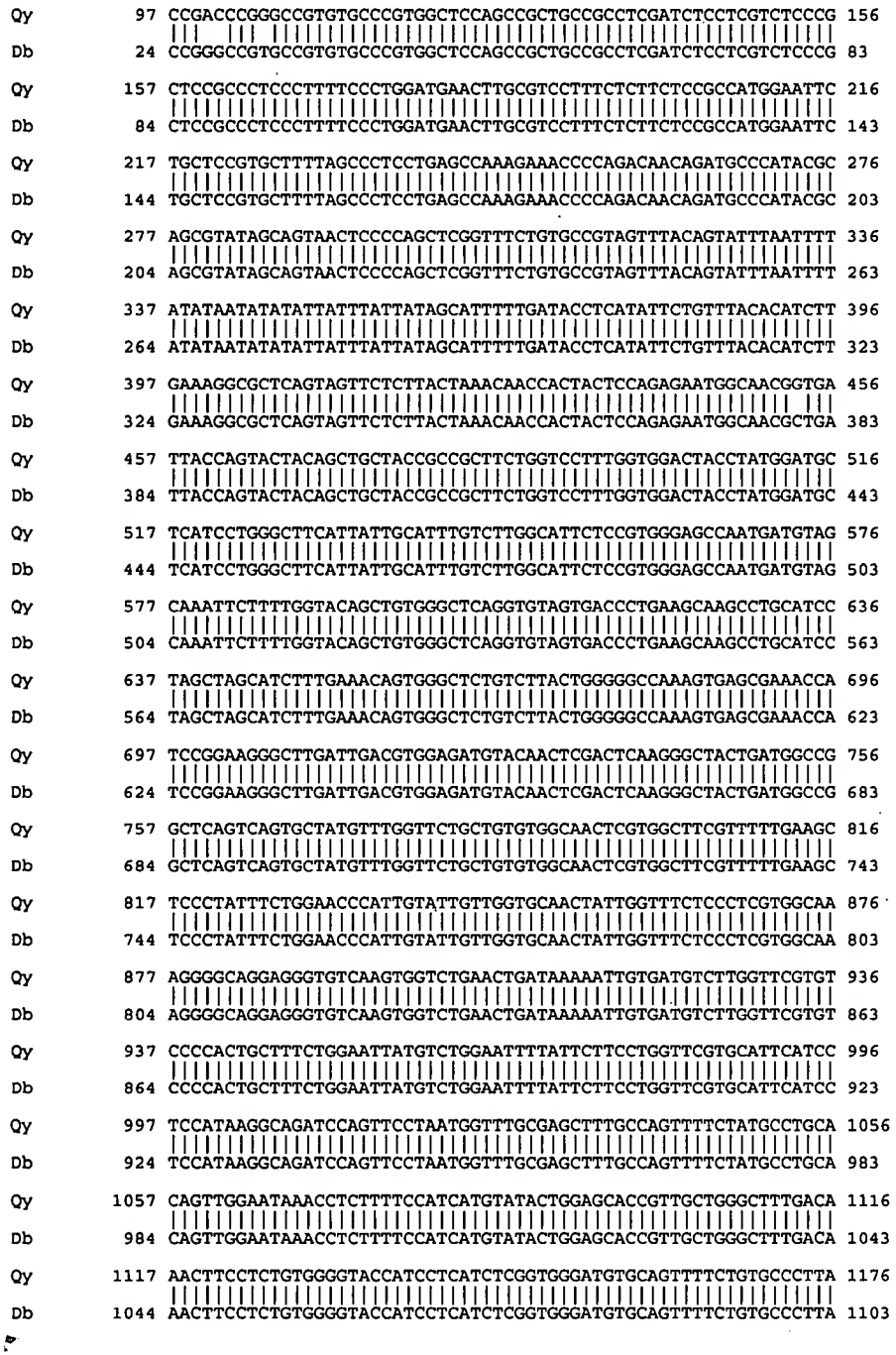
	16	73.6	2.2	4411529	3	US-09-103-840A-1	Sequence 1, Appli
	17	62.2	1.9	7218	1	US-08-232-463-14	Sequence 14, Appl
	18	58.6	1.8	1029	4	US-09-134-001C-2591	Sequence 2591, Ap
c	19	55.6	1.7	783	4	US-09-252-991A-6003	Sequence 6003, Ap
c	20	54.4	1.7	585	4	US-09-252-991A-6083	Sequence 6083, Ap
c	21	52.2	1.6	7218	1	US-08-232-463-14	Sequence 14, Appl
c	22	42.2	1.3	289	3	US-09-007-005-17	Sequence 17, Appl
c	23	42.2	1.3	289	3	US-09-244-796-17	Sequence 17, Appl
c	24	41.2	1.3	4403765	3	US-09-103-840A-2	Sequence 2, Appli
c	25	41.2	1.3	4411529	3	US-09-103-840A-1	Sequence 1, Appli
c	26	40.4	1.2	569	4	US-09-461-325-44	Sequence 44, Appl
c	27	40.4	1.2	31491	3	US-09-360-186-1	Sequence 1, Appli
c	28	40.2	1.2	1280	3	US-09-060-756-4	Sequence 4, Appli
c	29	40.2	1.2	1280	4	US-09-670-314-4	Sequence 4, Appli
	30	39.8	1.2	1410	4	US-09-634-238-191	Sequence 191, App
c	31	39.4	1.2	8257	4	US-09-595-684B-30	Sequence 30, Appl
c	32	39.4	1.2	8503	4	US-09-620-312D-130	Sequence 130, App
	33	39.4	1.2	16442	3	US-08-781-891-208	Sequence 208, App
	34	39.4	1.2	16442	4	US-09-618-166-208	Sequence 208, App
	35	39.2	1.2	521	3	US-09-488-744A-10	Sequence 10, Appl
c	36	39	1.2	1028	3	US-08-118-200-1	Sequence 1, Appli
c	37	39	1.2	1028	3	US-08-458-745-1	Sequence 1, Appli
c	38	38.8	1.2	3999	4	US-09-351-224E-9	Sequence 9, Appli
c	39	38.8	1.2	3999	4	US-09-677-488A-9	Sequence 9, Appli
c	40	38.8	1.2	3999	4	US-09-677-682B-9	Sequence 9, Appli
c	41	38.6	1.2	401	3	US-09-221-298-45	Sequence 45, Appl
c	42	37	1.1	366	4	US-09-389-681-278	Sequence 278, App
c	43	37	1.1	366	4	US-09-620-405B-278	Sequence 278, App
c	44	37	1.1	366	4	US-09-339-338-278	Sequence 278, App
c	45	37	1.1	366	4	US-09-433-826B-278	Sequence 278, App

ALIGNMENTS

RESULT 1
US-07-674-287B-1
; Sequence 1, Application US/07674287B
; Patent No. 5414076
; GENERAL INFORMATION:
; APPLICANT: Bryan Mark O'Hara
; TITLE OF INVENTION: Gibbon Ape Leukemia
; TITLE OF INVENTION: Virus Receptor
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Dr. Karen A. Lowney
; ADDRESSEE: American Cyanamid Company
; STREET: 1937 West Main Street
; STREET: P.O. Box 60
; CITY: Stamford
; STATE: CT
; COUNTRY: USA
; ZIP: 06904-0060
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy Disk
; COMPUTER: IBM PC AT
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: ASCII converted from IBM DW4
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/674,287B
; FILING DATE: 19910325
; CLASSIFICATION: 530
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Lowney, Karen A., Dr.
; REGISTRATION NUMBER: 31,274
; REFERENCE/DOCKET NUMBER: 31,104-01
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 203 321 2361
; TELEFAX: 203 321 2971
; TELEX: 710 474 4059
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 3211 Base Pairs
; TYPE: NUCLEOTIDE SEQUENCE
; STRANDEDNESS: Single
; TOPOLOGY: Linear
; MOLECULE TYPE: DNA
US-07-674-287B-1

Query Match 96.4%; Score 3172; DB 1; Length 3211;
Best Local Similarity 99.7%; Pred. No. 0;
Matches 3180; Conservative 0; Mismatches 8; Indels 1; Gaps 1;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RESULT 2
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; Sequence 1, Application US/08436900A
; Patent No. 5874264
;  GENERAL INFORMATION:
;    APPLICANT:  O'Hara, Bryan M.
;    TITLE OF INVENTION:  Gibbon Ape Leukemia Virus Receptors
;    NUMBER OF SEQUENCES:  4
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  American Home Products
;      STREET:  One Campus Drive
;      CITY:  Parsippany
;      STATE:  New Jersey
;      COUNTRY:  U.S.A.
;      ZIP:  07054
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/436,900A
;      FILING DATE:  08-MAY-1995
;      CLASSIFICATION:  536
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Barnhard, Elizabeth M.
;      REGISTRATION NUMBER:  31,088
;      REFERENCE/DOCKET NUMBER:  31,104-03
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  201-683-2158
;      TELEFAX:  201-683-4117
;  INFORMATION FOR SEQ ID NO:  1:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3211 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
US-08-436-900A-1

  Query Match           96.4%;  Score 3172;  DB 2;  Length 3211;
  Best Local Similarity   99.7%;  Pred. No. 0;
  Matches 3180;  Conservative    0;  Mismatches    8;  Indels    1;  Gaps    1;

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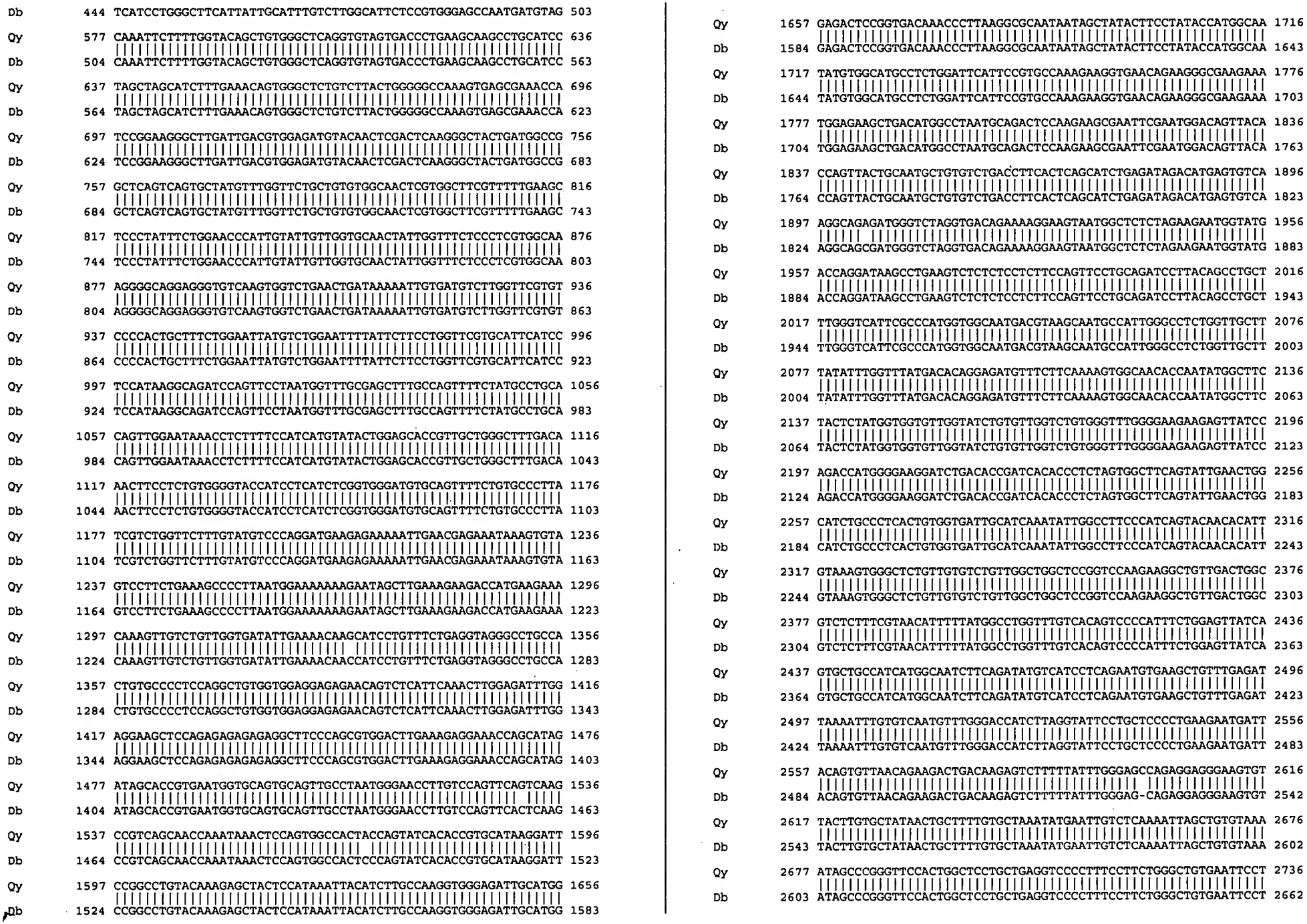
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Db 2184 CATCTGCCCTCACTGTGGTGATTGCATCAAATATTGGCCTTCCCATCAGTACAACACATT 2243
Qy 2317 GTAAAGTGGGCTCTGTTGTGTCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGC 2376
Db 2244 GTAAAGTGGGCTCTGTTGTGTCTGTTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGC 2303
Qy 2377 GTCTCTTTCGTAACATTTTTATGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCA 2436
Db 2304 GTCTCTTTCGTAACATTTTTATGGCCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATCA 2363
Qy 2437 GTGCTGCCATCATGGCAATCTTCAGATATGTCATCCTCAGAATGTGAAGCTGTTTGAGAT 2496
Db 2364 GTGCTGCCATCATGGCAATCTTCAGATATGTCATCCTCAGAATGTGAAGCTGTTTGAGAT 2423
Qy 2497 TAAAATTTGTGTCAATGTTTGGGACCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATT 2556
Db 2424 TAAAATTTGTGTCAATGTTTGGGACCATCTTAGGTATTCCTGCTCCCCTGAAGAATGATT 2483
Qy 2557 ACAGTGTTAACAGAAGACTGACAAGAGTCTTTTTATTTGGGAGCCAGAGGAGGGAAGTGT 2616
Db 2484 ACAGTGTTAACAGAAGACTGACAAGAGTCTTTTTATTTGGGAG-CAGAGGAGGGAAGTGT 2542
Qy 2617 TACTTGTGCTATAACTGCTTTTGTGCTAAATATGAATTGTCTCAAAATTAGCTGTGTAAA 2676
Db 2543 TACTTGTGCTATAACTGCTTTTGTGCTAAATATGAATTGTCTCAAAATTAGCTGTGTAAA 2602
Qy 2677 ATAGCCCGGGTTCCACTGGCTCCTGCTGAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCT 2736
Db 2603 ATAGCCCGGGTTCCACTGGCTCCTGCTGAGGTCCCCTTTCCTTCTGGGCTGTGAATTCCT 2662

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Qy       2737 GTACATATTTCTCTACTTTTTGTATCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTC 2796
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2663 GTACATATTTCTCTACTTTTTGTATCAGGCTTCAATTCCATTATGTTTTAATGTTGTCTC 2722

Qy       2797 TGAAGATGACTTGTGATTTTTTTTTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAG 2856
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2723 TGAAGATGACTTGTGATTTTTTTTTCTTTTTTTTAAACCATGAAGAGCCGTTTGACAGAG 2782

Qy       2857 CATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAAC 2916
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2783 CATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCACAGGGATTTAACAAC 2842

Qy       2917 AAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTG 2976
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2843 AAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAGTCCTTGGTACTCTG 2902

Qy       2977 CCCTCCTGTCAGTAGTGGCAGGATCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAG 3036
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2903 CCCTCCTGTCAGTAGTGGCAGGATCTATTGGCATATTCGGGAGCTTCTTAGAGGGATGAG 2962

Qy       3037 GTTCTTTGAACACAGTGAAAATTTAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTG 3096
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       2963 GTTCTTTGAACACAGTGAAAATTTAAATTAGTAACTTTTTTGCAAGCAGTTTATTGACTG 3022

Qy       3097 TTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAA 3156
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Db       3023 TTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAATCTTGGTTATTTCTTTAA 3082

Qy       3157 GATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAAT 3216
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Db       3083 GATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAACTTTGGGCAAGTTAAAT 3142

Qy       3217 GGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTT 3276
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       3143 GGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATAAAAAAGCCTACAGTTTT 3202

Qy       3277 TAGAAAAAA 3285
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RESULT 3
US-07-674-287B-3
; Sequence 3, Application US/07674287B
; Patent No. 5414076
;  GENERAL INFORMATION:
;    APPLICANT:  Bryan Mark O'Hara
;    TITLE OF INVENTION:  Gibbon Ape Leukemia
;    TITLE OF INVENTION:  Virus Receptor
;    NUMBER OF SEQUENCES:  3
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  Dr. Karen A. Lowney
;      ADDRESSEE:  American Cyanamid Company
;      STREET:  1937 West Main Street
;      STREET:  P.O. Box 60
;      CITY:  Stamford
;      STATE:  CT
;      COUNTRY:  USA
;      ZIP:  06904-0060
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy Disk
;      COMPUTER:  IBM PC AT
;      OPERATING SYSTEM:  MS-DOS
;      SOFTWARE:  ASCII converted from IBM DW4
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/07/674,287B
;      FILING DATE:  19910325
;      CLASSIFICATION:  530
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:
;      FILING DATE:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Lowney, Karen A., Dr.
;      REGISTRATION NUMBER:  31,274
;      REFERENCE/DOCKET NUMBER:  31,104-01
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  203 321 2361
;      TELEFAX:  203 321 2971
;      TELEX:  710 474 4059
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3260 BPs   681 Amino Acid Residues
;      TYPE:  NUCLEOTIDE AND AMINO ACID SEQUENCES
;      STRANDEDNESS:  Single
;      TOPOLOGY:  Linear
;    MOLECULE TYPE:  DNA and Protein
US-07-674-287B-3

  Query Match            64.8%;  Score 2133;  DB 1;  Length 3260;
  Best Local Similarity  84.5%;  Pred. No. 0;
  Matches 2662;  Conservative    0;  Mismatches  420;  Indels   67;  Gaps   21;

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              | || | |   ||| ||||||||||||||||||||||  ||||||||||||||||||||
Db        157 CACTTCGTCCCCCCAGGATGAACTTGCGTCCTTTCTCTAATCCGCCATGGAATTCTGCTC 216

Qy        222 CGTGCTTTTAGCCCTCCTGAGCCAAAGAAACCCCAGACAACAGATGCCCATAC---GCAG 278
              |||||||||||||||| |||||||||||||||||||||||||| ||||| ||    ||||
Db        217 CGTGCTTTTAGCCCTCCAGAGCCAAAGAAACCCCAGACAACAGACGCCCAGACGCAGCAG 276

Qy        279 CGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTTAT 338
              |||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        277 CGTATAGCAGTAACTCCCCAGCTCGGTTTCCGTGCCGTAGTTTACAGTATTTAATTTTAT 336

Qy        339 ATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTTGA 398
              |||||||||| |||||||||||||||||||     |   |  |||| ||||||||||||  |
Db        337 ATAATATATACTATTTATTATAGCATTTT---GATACCTCATTCCGTTTACACATCTCAA 393

Qy        399 AAGGCGCTCAGTAGTTCTCTTA----CTAAACAACCACTACTCCAGAGAA---------- 444
              ||| |||| |||| ||||||||     |||| |||||||||  | ||||||
Db        394 AAGCCGCTTAGTAATTCTCTTATTATTTAAAGAACCACTACACTAGAGAATGGAATCTAC 453

Qy        445 --TGGCAACGGTGATTACCAGTACTACAGCTGCTACCGCCGCTTCTGGTCCTTTGGTGGA 502
                |||||||| | | ||  |     | |||| ||| ||  |  ||    ||   |  ||
Db        454 TGTGGCAACGATTACTAGTACCCTAGCTGCTGTTACTGCTTCCGCTCCACCGAAGTATGA 513

Qy        503 CTACCTATGGATGCTCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCTCCGTGGG 562
              | | |||||||||||||||||||||||| ||||||||||||||||||||||||||||||
Db        514 CAATCTATGGATGCTCATCCTGGGCTTCATCATTGCATTTGTCTTGGCATTCTCCGTGGG 573

Qy        563 AGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAA 622
              |||||||||||||||||||| || |||||||||| |||||||||||||||||||||||||
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Qy        623 GCAAGCCTGCATCCTAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAA 682
              |||||||||||| |||||||||||| ||||| |||||||| | ||| ||||||||||||
Db        634 GCAAGCCTGCATCTTAGCTAGCATCTTCGAAACTGTGGGCTCCGCCTTGCTGGGGGCCAA 693

Qy        683 AGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGG 742
              |||||||||||||||||||| |||||||| || |||||| ||||||||    |||||||
Db        694 AGTGAGCGAAACCATCCGGAACGGCTTGATAGATGTGGAGCTGTACAACGAAACTCAAGA 753

Qy        743 GCTACTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGC 802
              || || ||||| ||||| ||||||||||||||||||||||||||||||| ||||||||
Db        754 TCTGCTCATGGCTGGCTCCGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAGCTCGTGGC 813

Qy        803 TTCGTTTTTGAAGCTCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTT 862
              |||||||||||||| || |||||||| |||||||||||||| ||||||| ||||||||
Db        814 TTCGTTTTTGAAGCTTCCGATTTCTGGGACCCATTGTATTGTCGGTGCAACCATTGGTTT 873

Qy        863 CTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGAT 922
              |||||| |||||||| |||||| ||||||||||||||||||||||||||||||||||||
Db        874 CTCCCTTGTGGCAAATGGGCAGAAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGAT 933

Qy        923 GTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGT 982
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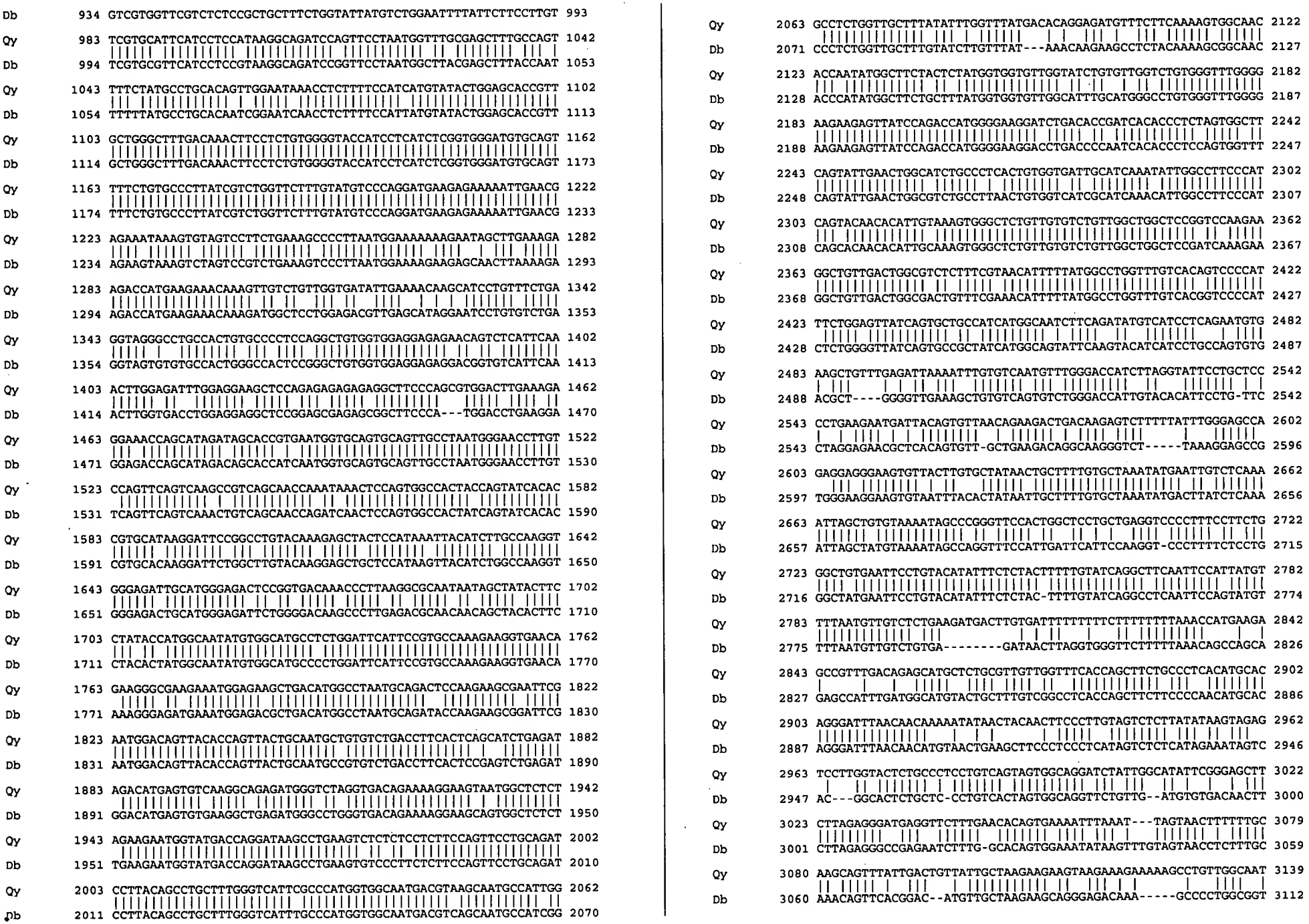
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Qy 1103 GCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGT 1162
Db 1114 GCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGT 1173
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Db 1174 TTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAATTGAACG 1233
Qy 1223 AGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGA 1282
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Qy 1283 AGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGA 1342
Db 1294 AGACCATGAAGAAACAAAGATGGCTCCTGGAGACGTTGAGCATAGGAATCCTGTGTCTGA 1353
Qy 1343 GGTAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCTCATTCAA 1402
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Qy 1403 ACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACTTGAAAGA 1462
Db 1414 ACTTGGTGACCTGGAGGAGGCTCCGGAGCGAGAGCGGCTTCCCA---TGGACCTGAAGGA 1470
Qy 1463 GGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGAACCTTGT 1522
Db 1471 GGAGACCAGCATAGACAGCACCATCAATGGTGCAGTGCAGTTGCCTAATGGGAACCTTGT 1530
Qy 1523 CCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTGGCCACTACCAGTATCACAC 1582
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Db 1591 CGTGCACAAGGATTCTGGCTTGTACAAGGAGCTGCTCCATAAGTTACATCTGGCCAAGGT 1650
Qy 1643 GGGAGATTGCATGGGAGACTCCGGTGACAAACCCTTAAGGCGCAATAATAGCTATACTTC 1702
Db 1651 GGGAGACTGCATGGGAGATTCTGGGGACAAGCCCTTGAGACGCAACAACAGCTACACTTC 1710
Qy 1703 CTATACCATGGCAATATGTGGCATGCCTCTGGATTCATTCCGTGCCAAAGAAGGTGAACA 1762
Db 1711 CTACACTATGGCAATATGTGGCATGCCCCTGGATTCATTCCGTGCCAAAGAAGGTGAACA 1770
Qy 1763 GAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATGCAGACTCCAAGAAGCGAATTCG 1822
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Qy 1823 AATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACCTTCACTCAGCATCTGAGAT 1882
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Qy 1883 AGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTGACAGAAAAGGAAGTAATGGCTCTCT 1942
Db 1891 GGACATGAGTGTGAAGGCTGAGATGGGCCTGGGTGACAGAAAAGGAAGCAGTGGCTCTCT 1950
Qy 1943 AGAAGAATGGTATGACCAGGATAAGCCTGAAGTCTCTCTCCTCTTCCAGTTCCTGCAGAT 2002
Db 1951 TGAAGAATGGTATGACCAGGATAAGCCTGAAGTGTCCCTTCTCTTCCAGTTCCTGCAGAT 2010
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Db 2011 CCTTACAGCCTGCTTTGGGTCATTTGCCCATGGTGGCAATGACGTCAGCAATGCCATCGG 2070
Qy 2063 GCCTCTGGTTGCTTTATATTTGGTTTATGACACAGGAGATGTTTCTTCAAAAGTGGCAAC 2122
Db 2071 CCCTCTGGTTGCTTTGTATCTTGTTTAT---AAACAAGAAGCCTCTACAAAAGCGGCAAC 2127
Qy 2123 ACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATCTGTGTTGGTCTGTGGGTTTGGGG 2182
Db 2128 ACCCATATGGCTTCTGCTTTATGGTGGTGTTGGCATTTGCATGGGCCTGTGGGTTTGGGG 2187
Qy 2183 AAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACACCGATCACACCCTCTAGTGGCTT 2242
Db 2188 AAGAAGAGTTATCCAGACCATGGGGAAGGACCTGACCCCAATCACACCCTCCAGTGGTTT 2247
Qy 2243 CAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATTGCATCAAATATTGGCCTTCCCAT 2302
Db 2248 CAGTATTGAACTGGCGTCTGCCTTAACTGTGGTCATCGCATCAAACATTGGCCTTCCCAT 2307
Qy 2303 CAGTACAACACATTGTAAAGTGGGCTCTGTTGTGTCTGTTGGCTGGCTCCGGTCCAAGAA 2362
Db 2308 CAGCACAACACATTGCAAAGTGGGCTCTGTTGTGTCTGTTGGCTGGCTCCGATCAAAGAA 2367
Qy 2363 GGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTATGGCCTGGTTTGTCACAGTCCCCAT 2422
Db 2368 GGCTGTTGACTGGCGACTGTTTCGAAACATTTTTATGGCCTGGTTTGTCACGGTCCCCAT 2427
Qy 2423 TTCTGGAGTTATCAGTGCTGCCATCATGGCAATCTTCAGATATGTCATCCTCAGAATGTG 2482
Db 2428 CTCTGGGGTTATCAGTGCCGCTATCATGGCAGTATTCAAGTACATCATCCTGCCAGTGTG 2487
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Db 2488 ACGCT----GGGGTTGAAAGCTGTGTCAGTGTCTGGGACCATTGTACACATTCCTG-TTC 2542
Qy 2543 CCTGAAGAATGATTACAGTGTTAACAGAAGACTGACAAGAGTCTTTTTATTTGGGAGCCA 2602
Db 2543 CTAGGAGAACGCTCACAGTGTT-GCTGAAGACAGGCAAGGGTCT-----TAAAGGAGCCG 2596
Qy 2603 GAGGAGGGAAGTGTTACTTGTGCTATAACTGCTTTTGTGCTAAATATGAATTGTCTCAAA 2662
Db 2597 TGGGAAGGAAGTGTAATTTACACTATAATTGCTTTTGTGCTAAATATGACTTATCTCAAA 2656
Qy 2663 ATTAGCTGTGTAAAATAGCCCGGGTTCCACTGGCTCCTGCTGAGGTCCCCTTTCCTTCTG 2722
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Qy 2723 GGCTGTGAATTCCTGTACATATTTCTCTACTTTTTGTATCAGGCTTCAATTCCATTATGT 2782
Db 2716 GGCTATGAATTCCTGTACATATTTCTCTAC-TTTTGTATCAGGCCTCAATTCCAGTATGT 2774
Qy 2783 TTTAATGTTGTCTCTGAAGATGACTTGTGATTTTTTTTTCTTTTTTTTAAACCATGAAGA 2842
Db 2775 TTTAATGTTGTCTGTGA--------GATAACTTAGGTGGGTTCTTTTTAAACAGCCAGCA 2826
Qy 2843 GCCGTTTGACAGAGCATGCTCTGCGTTGTTGGTTTCACCAGCTTCTGCCCTCACATGCAC 2902
Db 2827 GAGCCATTTGATGGCATGTACTGCTTTGTCGGCCTCACCAGCTTCTTCCCCAACATGCAC 2886
Qy 2903 AGGGATTTAACAACAAAAATATAACTACAACTTCCCTTGTAGTCTCTTATATAAGTAGAG 2962
Db 2887 AGGGATTTAACAACATGTAACTGAAGCTTCCCTCCCTCATAGTCTCTCATAGAAATAGTC 2946
Qy 2963 TCCTTGGTACTCTGCCCTCCTGTCAGTAGTGGCAGGATCTATTGGCATATTCGGGAGCTT 3022
Db 2947 AC---GGCACTCTGCTC-CCTGTCACTAGTGGCAGGTTCTGTTG--ATGTGTGACAACTT 3000
Qy 3023 CTTAGAGGGATGAGGTTCTTTGAACACAGTGAAAATTTAAAT---TAGTAACTTTTTTGC 3079
Db 3001 CTTAGAGGGCCGAGAATCTTTG-GCACAGTGGAAATATAAGTTTGTAGTAACCTCTTTGC 3059
Qy 3080 AAGCAGTTTATTGACTGTTATTGCTAAGAAGAAGTAAGAAAGAAAAAGCCTGTTGGCAAT 3139
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Qy       3140 CTTGGTTATTTCTTTAAGATTTCTGGCAGTGTGGGATGGATGAATGAAGTGGAATGTGAA 3199
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Db       3113 TGTGGTTATTCTTCTGAGATTTCTGGCAGTGTGGGATGGGTGAATGAAGTGGAATGTGAA 3172

Qy       3200 CTTTGGGCAAGTTAAATGGGACAGCCTTCCATGTTCATTTGTCTACCTCTTAACTGAATA 3259
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Db       3173 CTTTGGGCAAATTCAATGGGACAGCCTTCCATGTTCATCTGTCTACCTCTTAACTGAAT- 3231

Qy       3260 AAAAAGCCTACAGTTTTTAGAAAAAAAAA 3288
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Db       3232 AAAAAGCCTACAGTTTTTAAAAAAAAAAA 3260


RESULT 4
US-08-436-900A-3
; Sequence 3, Application US/08436900A
; Patent No. 5874264
;  GENERAL INFORMATION:
;    APPLICANT:  O'Hara, Bryan M.
;    TITLE OF INVENTION:  Gibbon Ape Leukemia Virus Receptors
;    NUMBER OF SEQUENCES:  4
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  American Home Products
;      STREET:  One Campus Drive
;      CITY:  Parsippany
;      STATE:  New Jersey
;      COUNTRY:  U.S.A.
;      ZIP:  07054
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  PatentIn Release #1.0, Version #1.30
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/436,900A
;      FILING DATE:  08-MAY-1995
;      CLASSIFICATION:  536
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Barnhard, Elizabeth M.
;      REGISTRATION NUMBER:  31,088
;      REFERENCE/DOCKET NUMBER:  31,104-03
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  201-683-2158
;      TELEFAX:  201-683-4117
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3260 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  DNA (genomic)
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  443..2488
US-08-436-900A-3

  Query Match             64.8%;  Score 2131.4;  DB 2;  Length 3260;
  Best Local Similarity   84.5%;  Pred. No. 0;
  Matches 2661;  Conservative    0;  Mismatches  421;  Indels   67;  Gaps   21;

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              | || | |   ||| ||||||||||||||||||||||  ||||||||||||||||||||
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Qy        279 CGTATAGCAGTAACTCCCCAGCTCGGTTTCTGTGCCGTAGTTTACAGTATTTAATTTTAT 338
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Db        277 CGTATAGCAGTAACTCCCCAGCTCGGTTTCCGTGCCGTAGTTTACAGTATTTAATTTTAT 336

Qy        339 ATAATATATATTATTTATTATAGCATTTTTGATACCTCATATTCTGTTTACACATCTTGA 398
              |||||||||| |||||||||||||||||    |  |  |||| ||||||||||||  |
Db        337 ATAATATATACTATTTATTATAGCATTTT---GATACCTCATTCCGTTTACACATCTCAA 393

Qy        399 AAGGCGCTCAGTAGTTCTCTTA----CTAAACAACCACTACTCCAGAGAA---------- 444
              ||| |||| |||| ||||||||     |||| |||||||| | ||||||
Db        394 AAGCCGCTTAGTAATTCTCTTATTATTTAAAGAACCACTACACTAGAGAATGGAATCTAC 453

Qy        445 --TGGCAACGGTGATTACCAGTACTACAGCTGCTACCGCCGCTTCTGGTCCTTTGGTGGA 502
                |||||||| | | ||  |       | |||| ||| ||  |  ||   ||   |  ||
Db        454 TGTGGCAACGATTACTAGTACCCTAGCTGCTGTTACTGCTTCCGCTCCACCGAAGTATGA 513

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Db        514 CAATCTATGGATGCTCATCCTGGGCTTCATCATTGCATTTGTCTTGGCATTCTCCGTGGG 573

Qy        563 AGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAA 622
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Db        574 AGCCAATGATGTAGCAAATTCGTTCGGTACAGCTGTAGGCTCAGGTGTAGTGACCCTGAA 633

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Db        634 GCAAGCCTGCATCTTAGCTAGCATCTTCGAAACTGTGGGCTCCGCCTTGCTGGGGGCCAA 693

Qy        683 AGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGG 742
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Db        694 AGTGAGCGAAACCATCCGGAACGGCTTGATAGATGTGGAGCTGTACAACGAAACTCAAGA 753

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Db        754 TCTGCTCATGGCTGGCTCCGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAGCTCGTGGC 813

Qy        803 TTCGTTTTTGAAGCTCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTT 862
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Db        814 TTCGTTTTTGAAGCTTCCGATTTCTGGGACCCATTGTATTGTCGGTGCAACCATTGGTTT 873

Qy        863 CTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGAT 922
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Db        874 CTCCCTTGTGGCAAATGGGCAGAAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGAT 933

Qy        923 GTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGT 982
              ||| |||||||| || || |||||||||| |||||||||||||||||||||||||| ||
Db        934 GTCGTGGTTCGTCTCTCCGCTGCTTTCTGGTATTATGTCTGGAATTTTATTCTTCCTTGT 993

Qy        983 TCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGT 1042
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Qy       1043 TTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAGCACCGTT 1102
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Db       1054 TTTTTATGCCTGCACAATCGGAATCAACCTCTTTTCCATTATGTATACTGGAGCACCGTT 1113

Qy       1103 GCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGT 1162
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db       1114 GCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGATGTGCAGT 1173

Qy       1163 TTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAATTGAACG 1222
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Qy       1223 AGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCTTGAAAGA 1282
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Qy       1283 AGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTGTTTCTGA 1342
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              |||||  |||||||| ||| |||| |||||||||||||||||| || || ||||||||
Db       1354 GGTAGTGTGTGCCACTGGGCCACTCCGGGCTGTGGTGGAGGAGAGGACGGTGTCATTCAA 1413

Qy       1403 ACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACTTGAAAGA 1462
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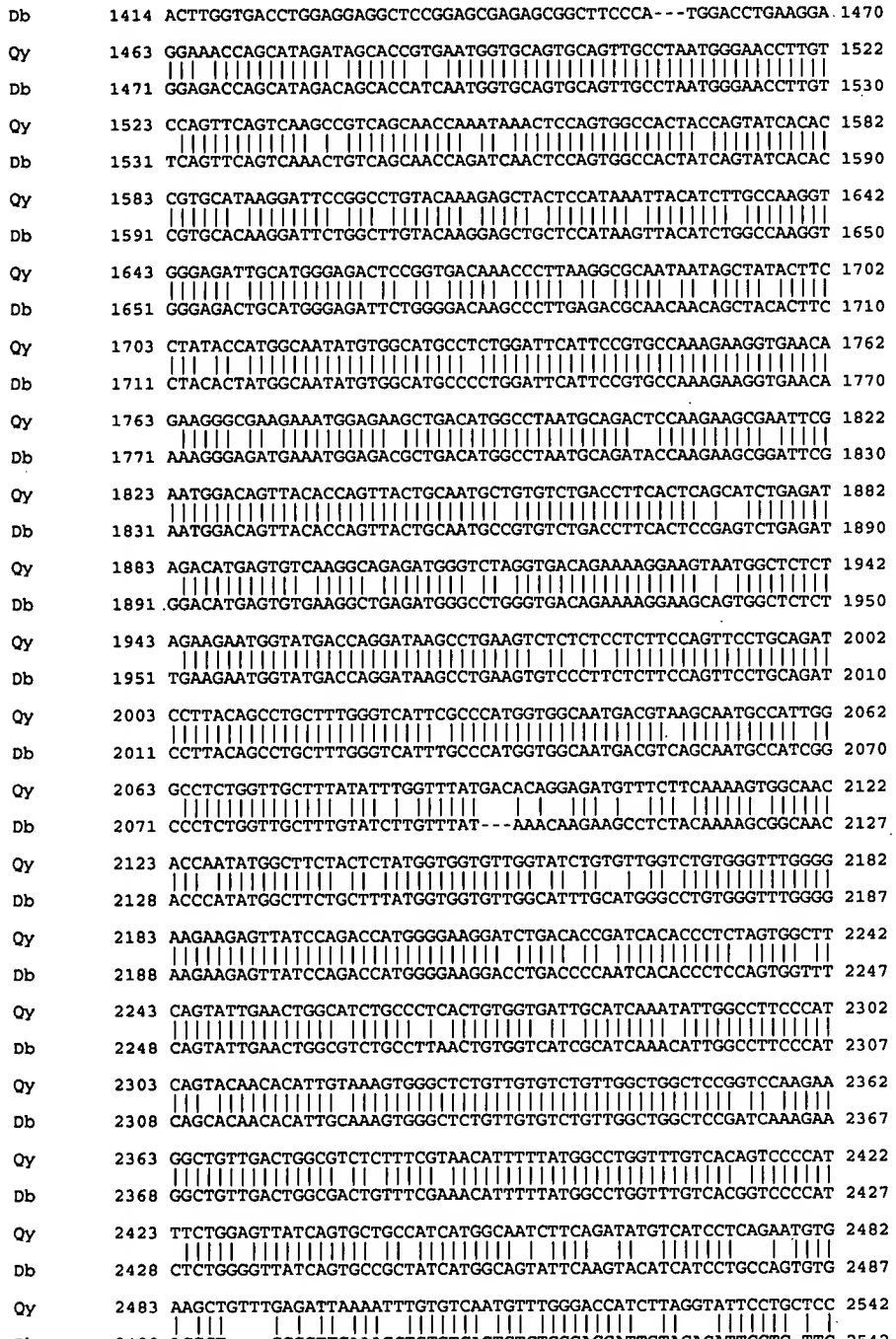

RESULT 5
US-08-050-684-1
; Sequence 1, Application US/08050684
; Patent No. 5550221
; GENERAL INFORMATION:
; APPLICANT: Johann Dr., Stephen V.
; APPLICANT: Van Zeijl Dr., Marja
; APPLICANT: O'Hara Dr., Bryan M.
; TITLE OF INVENTION: Amphotropic Virus Receptor
; NUMBER OF SEQUENCES: 2
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: American Cyanamid Company
; STREET: 1937 West Main Street
; CITY: Stamford
; STATE: CT
; COUNTRY: United States of America
; ZIP: 06904-0060
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk

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;       COMPUTER:  IBM PC compatible
;       OPERATING SYSTEM:  PC-DOS/MS-DOS
;       SOFTWARE:  PatentIn Release #1.0, Version #1.25
;     CURRENT APPLICATION DATA:
;       APPLICATION NUMBER:  US/08/050,684
;       FILING DATE:  16-APR-1993
;       CLASSIFICATION:  536
;     ATTORNEY/AGENT INFORMATION:
;       NAME:  Lowney Dr., Karen A
;       REGISTRATION NUMBER:  31,274
;       REFERENCE/DOCKET NUMBER:  31937-00
;     TELECOMMUNICATION INFORMATION:
;       TELEPHONE:  203-321-2361
;       TELEFAX:  203-321-2971
;       TELEX:  710-474-4059
;   INFORMATION FOR SEQ ID NO:  1:
;     SEQUENCE CHARACTERISTICS:
;       LENGTH:  3175 base pairs
;       TYPE:  nucleic acid
;       STRANDEDNESS:  single
;       TOPOLOGY:  linear
;     MOLECULE TYPE:  cDNA
;     HYPOTHETICAL:  NO
;     ANTI-SENSE:  NO
;     FEATURE:
;       NAME/KEY:  CDS
;       LOCATION:  244..2202
US-08-050-684-1

  Query Match             18.8%;  Score 617.2;  DB 1;  Length 3175;
  Best Local Similarity   60.3%;  Pred. No. 7.5e-169;
  Matches 1203;  Conservative    0;  Mismatches  718;  Indels   75;  Gaps    8;

Qy        496 TGGTGGACTACCTATGGATGCTCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCT 555
Db        251 TGGATGAGTATTTGTGGATGGTCATTTTGGGTTTCATCATAGCTTTCATCTTGGCCTTTT 310

Qy        556 CCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGA 615
Db        311 CTGTTGGTGCAAACGATGTTGCCAACTCCTTTGGTACAGCCGTGGGCTCTGGTGTGGTGA 370

Qy        616 CCCTGAAGCAAGCCTGCATCCTAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGG 675
Db        371 CCTTGAGGCAGGCATGCATTTTAGCTTCAATATTTGAAACCACCGGCTCCGTGTTACTAG 430

Qy        676 GGGCCAAAGTGAGCGAAACCATCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGA 735
Db        431 GCGCCAAAGTAGGAGAAACCATTCGCAAAGGTATCATTGACGTGAACCTGTACAACGAGA 490

Qy        736 CTCAAGGGCTACTGATGGCCGGCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAAC 795
Db        491 CGGTGGAGACTCTCATGGCTGGGGAAGTTAGTGCCATGGTTGGTTCCGCTGTGTGGCAGC 550

Qy        796 TCGTGGCTTCGTTTTGAAGCTCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTA 855
Db        551 TGATTGCTTCCTTCCTGAGGCTTCCAATCTCAGGAACGCACTGCATTGTGGGTTCTACTA 610

Qy        856 TTGGTTTCTCCCTCGTGGCAAAGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAA 915
Db        611 TAGGATTCTCACTGGTCGCAATCGGTACCAAAGGTGTGCAGTGGATGGAGCTTGTCAAGA 670

Qy        916 TTGTGATGTCTTGGTTCGTGTCCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCT 975
Db        671 TTGTTGCTTCTTGGTTTATATCTCCACTGTTGTCTGGTTTCATGTCTGGCCTGCTGTTTG 730

Qy        976 TCCTGGTTCGTGCATTCATCCTCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTT 1035
Db        731 TACTCATCAGAATTTTCATCTTAAAAAAGGAAGACCCTGTTCCCAATGGCCTCCGGGCAC 790

Qy       1036 TGCCAGTTTTCTATGCCTGCACAGTTGGAATAAACCTCTTTTCCATCATGTATACTGGAG 1095
Db        791 TCCCAGTATTCTATGCTGCTACCATAGCAATCAATGTCTTTTCCATCATGTACACAGGAG 850

Qy       1096 CACCGTTGCTGGGCTTTGACAAACTTCCTCTGTGGGGTACCATCCTCATCTCGGTGGGAT 1155
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Qy       1156 GTGCAGTTTTCTGTGCCCTTATCGTCTGGTTCTTTGTATGTCCCAGGATGAAGAGAAAAA 1215
Db        908 TCGCCCTCCTGTTCGCTTTTTTGTGTGGCTCTTCGTGTGTCCGTGGATGCGGAGGAAAA 967

Qy       1216 TTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGCCCCTTAATGGAAAAAAAGAATAGCT 1275
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Qy       1276 TGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTTGGTGATATTGAAAACAAGCATCCTG 1335
Db       1028 AGGTTCAGGAAGCAGAGTCCCCAGTATT-----------------TAAAGAGCTACCAGG 1070

Qy       1336 TTTCTGAGGTAGGGCCTGCCACTGTGCCCCTCCAGGCTGTGGTGGAGGAGAGAACAGTCT 1395
Db       1071 TGCCAAGGCTAATGATGACAGCACCATCCCGCTCACGGGAGCAGCAGGGGAG-------- 1122

Qy       1396 CATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAGAGAGAGAGGCTTCCCAGCGTGGACT 1455
Db       1123 -----ACACTGGGGACCTCGGAAGGCACTTCTGCG-------GGCAGCCACCCTCGGGCT 1170

Qy       1456 TGAAAGAGGAAACCAGCATAGATAGCACCGTGAATGGTGCAGTGCAGTTGCCTAATGGGA 1515
Db       1171 GCATACGGAAGAGCACTGTCCATGACCCATGGCTCTGTGAAATCGCCCATCTCCAACGGC 1230

Qy       1516 ACCTTGTCCAGTTCAGTCAAGCCGTCAGCAACCAAATAAACTCCAGTGGCCACTACCAGT 1575
Db       1231 ACCTT-----------------CGGCTTCGACGGCCACACCAGGAGCGACGGTCATGTGT 1273

Qy       1576 ATCACACCGTGCATAAGGATTCCGGCCTGTACAAAGAGCTACTCCATAAATTACATCTTG 1635
Db       1274 ACCACACCGTGCACAAAGACTCGGGGCTCTACAAAGATCTGCTGCACAAAATCCACATCG 1333

Qy       1636 CCAAGGTGGGAGATTGCATGGGAGACTCCGGTGACAA------ACCCTTAAGGCGCAATA 1689
Db       1334 ACAGGGGCCCCGAGGAGAAGCCAGCCCAGGAAAGCAACTACCGGCTGCTCCGCCGAAACA 1393

Qy       1690 ATAGCTATACTTCCTATACCATGGCAATATGTGGCATGCCTCTGGATTCATTCCGTGCCA 1749
Db       1394 ACAGTTACACCTGCTACACCGCAGCCATTTGTGGGCTGCCAGTGCACGCCACCTTTCGAG 1453

Qy       1750 AAGAAGGTGAACAGAAGGGCGAAGAAATGGAGAAGCTGACATGGCCTAATG---CAGACT 1806
Db       1454 CTGCGGACTCATCGGCCCCAGAGGACAGTGAGAAGCTGGTGGGCGACACCGTGTCCTACT 1513

Qy       1807 CCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACCTTC 1866
Db       1514 CCAAGAAGAGGCTGCGCTACGACAGCTACTCGAGCTACTGTAACGCGGTGGCAGAGGCGG 1573

Qy       1867 ACTCAGCATCTGAGA---------TAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTG 1917
Db       1574 AGATCGAGGCGGAGGAGGGCGGCGTGGAGATGAAGCTGGCGTCGGAGCTGGCCGACCCTG 1633

Qy       1918 ACAGAAAAGGAAGTAATGGCTCTCTAGAAGAATGGTATGACCAGGATAAGCCTGAAGTCT 1977
Db       1634 ACCAGCCGCGAGAGGACCCTGCAGAGGAGGAGAAGGAGGAGAAGGACGCACCCGAGGTTC 1693

Qy       1978 CTCTCCTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTG 2037
Db       1694 ACCTCCTGTTCCATTTCCTGCAGGTCCTCACCGCCTGTTTCGGGTCCTTTGCTCACGGCG 1753

Qy       2038 GCAATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAG 2097
Db       1754 GCAATGACGTGAGTAATGCCATCGGTCCCCTGGTAGCCTTGTGGCTGATTTACAAACAAG 1813

Qy       2098 GAGATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTA 2157
Db       1814 GCGGGGTAACGCAAGAAGCAGCTACACCCGTCTGGCTGCTGTTTTATGGAGGAGTTGGAA 1873

Qy       2158 TCTGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGA 2217
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Qy       2218 CACCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGA 2277

Db       1934 CTCCCATCACGCCGTCCAGCGGCTTCACGATCGAGCTGGCCTCAGCCTTCACAGTGGTGA 1993

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Db       1994 TCGCCTCCAACATCGGGCTTCCAGTCAGCACCACGCACTGTAAGGTGGGCTCGGTGGTGG 2053

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Qy       2458 TCAGATATGTCATCCT 2473

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RESULT 6
US-08-582-719-1
; Sequence 1, Application US/08582719
; Patent No. 5633348
; GENERAL INFORMATION:
; APPLICANT: Johann Dr., Stephen V.
; APPLICANT: Van Zeijl Dr., Marja
; APPLICANT: O'Hara Dr., Bryan M.
; TITLE OF INVENTION: Amphotropic Virus Receptor
; NUMBER OF SEQUENCES: 2
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: American Cyanamid Company
; STREET: 1937 West Main Street
; CITY: Stamford
; STATE: CT
; COUNTRY: United States of America
; ZIP: 06904-0060
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/582,719
; FILING DATE: 04-JAN-1996
; CLASSIFICATION: 530
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/050,684
; FILING DATE: 16-APR-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Lowney Dr., Karen A
; REGISTRATION NUMBER: 31,274
; REFERENCE/DOCKET NUMBER: 31937-00
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 203-321-2361
; TELEFAX: 203-321-2971
; TELEX: 710-474-4059
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 3175 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 244..2202

US-08-582-719-1

Query Match 18.8%; Score 617.2; DB 1; Length 3175;
Best Local Similarity 60.3%; Pred. No. 7.5e-169;
Matches 1203; Conservative 0; Mismatches 718; Indels 75; Gaps 8;

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Qy        556 CCGTGGGAGCCAATGATGTAGCAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGA 615

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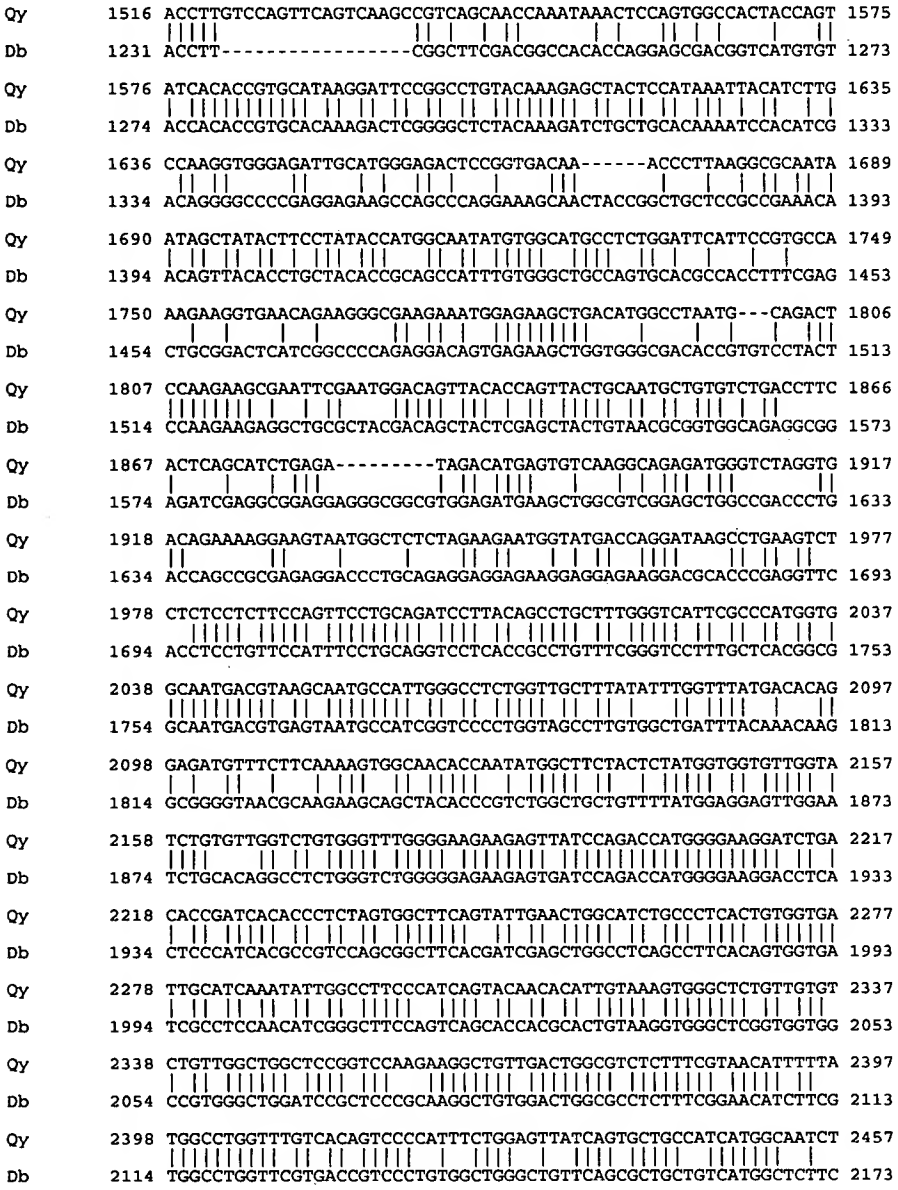
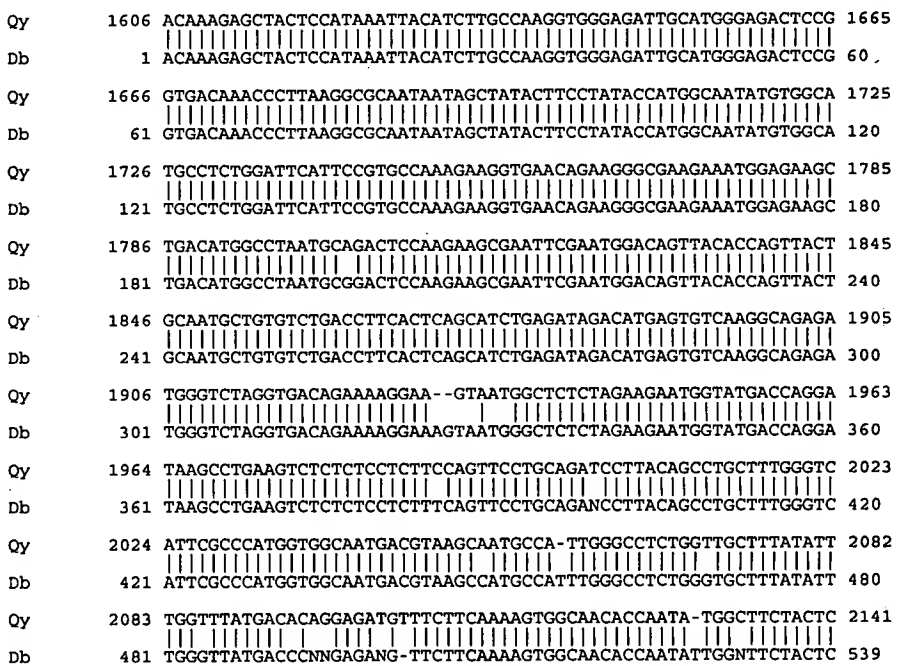


RESULT 7
US-09-328-111-623
; Sequence 623, Application US/09328111

; Patent No. 6262333
; GENERAL INFORMATION:
; APPLICANT: Endege, Wilson O.
; APPLICANT: Steinmann, Kathleen E.
; APPLICANT: Astle, Jon H.
; APPLICANT: Burgess, Christopher C.
; APPLICANT: Bushnell, Steven E.
; APPLICANT: Carroll III, Eddie
; APPLICANT: Catino, Theodore J.
; APPLICANT: Derti, Adnan
; APPLICANT: Ford, Donna M.
; APPLICANT: Lewis, Marcia E.
; APPLICANT: Monahan, John E.
; APPLICANT: Schlegel, Robert
; TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
; TITLE OF INVENTION: PRODUCTS
; FILE REFERENCE: CCD-257 (US)
; CURRENT APPLICATION NUMBER: US/09/328,111
; CURRENT FILING DATE: 1999-06-08
; EARLIER APPLICATION NUMBER: US 60/088,801
; EARLIER FILING DATE: 1998-06-10
; NUMBER OF SEQ ID NOS: 850
; SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 623
; LENGTH: 662
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(662)
; OTHER INFORMATION: n = A,T,C or G
US-09-328-111-623

Query Match 15.1%; Score 495.2; DB 3; Length 662;
Best Local Similarity 90.2%; Pred. No. 8e-134;
Matches 591; Conservative 0; Mismatches 54; Indels 10; Gaps 6;



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Qy       2142 TAT--GGTGGTGTTGGTATCTGTGTTGG---TCTGTGGGTTTGGGGAAGAAGAGTTATCC 2196
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Db        540 TANTGGNGGGGGTTGGGATCTGNGGTTGGTCTGTGGGGTTTGGGGAAAAAAAAGTTTTCC 599

Qy       2197 AGACCATGGGGAAGGATCTGACACCGATCACACCCTCTAGTGGCTTCAGTATTGA 2251
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RESULT 8
US-09-328-111-521
; Sequence 521, Application US/09328111
; Patent No. 6262333
; GENERAL INFORMATION:
;  APPLICANT: Endege, Wilson O.
;  APPLICANT: Steinmann, Kathleen E.
;  APPLICANT: Astle, Jon H.
;  APPLICANT: Burgess, Christopher C.
;  APPLICANT: Bushnell, Steven E.
;  APPLICANT: Carroll III, Eddie
;  APPLICANT: Catino, Theodore J.
;  APPLICANT: Derti, Adnan
;  APPLICANT: Ford, Donna M.
;  APPLICANT: Lewis, Marcia E.
;  APPLICANT: Monahan, John E.
;  APPLICANT: Schlegel, Robert
;  TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
;  TITLE OF INVENTION: PRODUCTS
;  FILE REFERENCE: CCD-257 (US)
;  CURRENT APPLICATION NUMBER: US/09/328,111
;  CURRENT FILING DATE: 1999-06-08
;  EARLIER APPLICATION NUMBER: US 60/088,801
;  EARLIER FILING DATE: 1998-06-10
;  NUMBER OF SEQ ID NOS: 850
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 521
;   LENGTH: 613
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(613)
;   OTHER INFORMATION: n = A,T,C or G
US-09-328-111-521

  Query Match             13.7%;  Score 451.4;  DB 3;  Length 613;
  Best Local Similarity   99.2%;  Pred. No. 4.2e-121;
  Matches  474;  Conservative    0;  Mismatches    2;  Indels    2;  Gaps    2;

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Qy       1191 GTATGTCCCAGGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGC 1250
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db         61 GTATGTCCCAGGATGAAGAGAAAAATTGAACGAGAAATAAAGTGTAGTCCTTCTGAAAGC 120

Qy       1251 CCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTT 1310
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        121 CCCTTAATGGAAAAAAAGAATAGCTTGAAAGAAGACCATGAAGAAACAAAGTTGTCTGTT 180

Qy       1311 GGTGATATTGAAAACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCACTGTGCCCCTCCAG 1370
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        181 GGTGATATTGAAAACAAGCATCCTGTTTCTGAGGTAGGGCCTGCCACTGTGCCCCTCCAG 240

Qy       1371 GCTGTGGTGGAGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAG 1430
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        241 GCTGTGGTGGAGGAGAGAACAGTCTCATTCAAACTTGGAGATTTGGAGGAAGCTCCAGAG 300

Qy       1431 AGAGAGAGGCTTCCCAGCGTGGACTTGAAAGAGGAAACCAGCATAGATAGCACCGTGAAT 1490
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        301 AGAGAGAGGCTTCCCAGCGTGGACTTGAAAGAGGAAACCAGCATAGATAGCACCGTGAAT 360

Qy       1491 GGTGCAGTGCAGTTGCCTAATGGGAACCTTGTCCAGTTCAGTC-AAGCCGTCAGCAACCA 1549
              ||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||
Db        361 GGTGCAGTGCAGTTGCCTAATGGGAACCTTGTCCAGTTCAGTCAAAGCCGTCAGCAACCA 420

Qy       1550 AATAAACTCCAGTGGCCACTACCAGTATCACACCGTGCAT-AAGGATTCCGGCCTGTA 1606
              |||||||| |||||||||||||||||||||||||||||| |||||||||||| |||||
Db        421 AATAAACTNCAGTGGCCACTACCAGTATCACACCGTGCATAAAGGATTCCGGGCTGTA 478


RESULT 9
US-09-328-111-269/c
; Sequence 269, Application US/09328111
; Patent No. 6262333
; GENERAL INFORMATION:
;  APPLICANT: Endege, Wilson O.
;  APPLICANT: Steinmann, Kathleen E.
;  APPLICANT: Astle, Jon H.
;  APPLICANT: Burgess, Christopher C.
;  APPLICANT: Bushnell, Steven E.
;  APPLICANT: Carroll III, Eddie
;  APPLICANT: Catino, Theodore J.
;  APPLICANT: Derti, Adnan
;  APPLICANT: Ford, Donna M.
;  APPLICANT: Lewis, Marcia E.
;  APPLICANT: Monahan, John E.
;  APPLICANT: Schlegel, Robert
;  TITLE OF INVENTION: NOVEL HUMAN GENES AND GENE EXPRESSION
;  TITLE OF INVENTION: PRODUCTS
;  FILE REFERENCE: CCD-257 (US)
;  CURRENT APPLICATION NUMBER: US/09/328,111
;  CURRENT FILING DATE: 1999-06-08
;  EARLIER APPLICATION NUMBER: US 60/088,801
;  EARLIER FILING DATE: 1998-06-10
;  NUMBER OF SEQ ID NOS: 850
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 269
;   LENGTH: 643
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(643)
;   OTHER INFORMATION: n = A,T,C or G
US-09-328-111-269

  Query Match             11.9%;  Score 393;  DB 3;  Length 643;
  Best Local Similarity   91.7%;  Pred. No. 4.2e-104;
  Matches  465;  Conservative    0;  Mismatches   35;  Indels    7;  Gaps    5;

Qy       1805 CTCCAAGAAGCGAATTCGAATGGACAGTTACACCAGTTACTGCAATGCTGTGTCTGACCT 1864
              |||||||||| || ||  ||||      | | ||||||||||||||||| ||||||| | |
Db        505 CTCCAAGAAGGGANTTNAAATG--NCAGTNCCCCAGTTACTGCAATGCNGTGTCTGNCTT 448

Qy       1865 TCACTCAGCATCTG-AGATAGACATGAGTGTCAAGGCAGAGATGGGTCTAGGTGACAGAA 1923
              ||| | ||| | |  |||||| ||| ||||  ||| || |||||||| |||||| |||||
Db        447 TCATTNAGCNTTTTAAGATAGCCATAAGTGCAAAGNCAAAGATGGGTTTAGGTGNCAGAA 388

Qy       1924 AAGGAAGTAATGGCTCT--CTAGAAGAATGGTATGACCAGGATAAG-CCTGAAGTCTCTC 1980
              ||| |||||| || |||      |||||||||| |||||||||| ||||||||||||
Db        387 AAGAAAGTAAGGGTTCTTCTAGAAAGAATGGTATGCCCAGGATAAGCCCTGAAGTCTCTC 328

Qy       1981 TCC-TCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTGGC 2039
              ||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        327 TCCNTCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTGGC 268

Qy       2040 AATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAGGA 2099
              |||||||||||||||||||||||||||||||||||||||||||||||||||| |||||||
Db        267 AATGACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGGCACAGGA 208

Qy       2100 GATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATC 2159
              ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        207 GATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATC 148
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Qy       2160 TGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACA 2219
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db        147 TGTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACA 88

Qy       2220 CCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATT 2279
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db          87 CCGATCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATT 28

Qy       2280 GCATCAAATATTGGCCTTCCCATCAGT 2306
             ||||||||||||||||||||||||||||
Db          27 GCATCAAATATTGGCCTTCCCATCAGT 1


RESULT 10
US-09-557-884-1
; Sequence 1, Application US/09557884
; Patent No. 6506581
;   GENERAL INFORMATION:
;        APPLICANT: Fleischmann et al.
;        TITLE OF INVENTION: The Nucleotide sequence of
;                            the Haemophilus influenzae Rd Genome, Fragments
;                            Thereof, and Uses Thereof
;        NUMBER OF SEQUENCES: 1
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: Human Genome Sciences, Inc.
;             STREET: 9410 Key West Avenue
;             CITY: Rockville
;             STATE: MD
;             COUNTRY: USA
;             ZIP: 20850
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: 3 1/2 inch diskette
;             COMPUTER: Dell Pentium
;             OPERATING SYSTEM: MS DOS v6.22
;             SOFTWARE: ASCII Text
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/557,884
;             FILING DATE: 25-Apr-2000
;             CLASSIFICATION: <Unknown>
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: 08/476,102
;             FILING DATE: JUN-5-1995
;        ATTORNEY/AGENT INFORMATION:
;             NAME: Michelle S. Marks
;             REGISTRATION NUMBER: 41,971
;             REFERENCE/DOCKET NUMBER: PB186P3
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: 301-309-8504
;             TELEFAX: 301-309-8439
;   INFORMATION FOR SEQ ID NO: 1:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 1830121 base pairs
;             TYPE: nucleic acid
;             STRANDEDNESS: double
;             TOPOLOGY: linear
;        SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-557-884-1

  Query Match             3.0%;  Score 99.8;  DB 4;  Length 1830121;
  Best Local Similarity   52.7%;  Pred. No. 1.9e-16;
  Matches  240;  Conservative    0;  Mismatches  212;  Indels    3;  Gaps    1;

Qy       1984 TCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTGGCAATG 2043
             |||||    |  |   | |  | || || ||       | |||| || || |||   ||||
Db    1670240 TCTTCAGTATTTTAATGTTATTAACCGCTTGTGCGATGGCATTTGCGCACGGTTCTAATG 1670299

Qy       2044 ACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTA---TATTTGGTTTATGACACAGGAG 2100
             |||||  |||||| ||||| |||||   |||  |    | | | ||  ||||    ||
Db    1670300 ACGTAGCCAATGCAATTGGCCCTCTTTCTGCAGTGGTCTCTATTGTAAATGAAGGCGGTA 1670359

Qy       2101 ATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATCT 2160
             |  ||  ||||   |  ||   | |    ||| || ||| ||  ||||   | |||||
Db    1670360 AAATTGTTTCAGGTGGAGCCTTAACTTGGTGGATTTTACCCTTAGGTGCATTAGGTATTG 1670419

Qy       2161 GTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACAC 2220
             ||| ||  |      |  |||| || ||| ||   ||| | ||    |   | ||
Db    1670420 CTGTGGGCTTAATTACTATGGGACAAAAAGTGATGGCGACTGTTGGATCTGGGATCACCG 1670479

Qy       2221 CGATCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATTG 2280
              | || ||     |||||||     | || | ||  ||||  | || || ||  | |
Db    1670480 ATTTAACGCCAAGCCGTGGCTTTGCCGCTCAATTTGCTACTGCGATGACCGTTGTCGTGG 1670539

Qy       2281 CATCAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGTCTG 2340
             |||||   | ||| | ||||||   ||||||||       || ||  || || |   |
Db    1670540 CATCAGGCACAGGCTTACCCATCTCAACAACACAAACACTTGTTGGAGCTATTTTAGGTA 1670599

Qy       2341 TTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTATGG 2400
             | ||||     ||    |     ||  |  | |       |  | || |||||  |||
Db    1670600 TCGGCTTTGCACGTGGTATCGCAGCACTGAATTTAACGGTTATCCGAAACATCATTAGTT 1670659

Qy       2401 CCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATC 2435
             | ||| |||||||| | ||      || | || ||
Db    1670660 CTTGGATTGTCACATTACCAGCAGGTGCATTTTTC 1670694


RESULT 11
US-09-643-990A-1
; Sequence 1, Application US/09643990A
; Patent No. 6528289
;   GENERAL INFORMATION:
;        APPLICANT: Robert D. Fleischmann
;                   Mark D. Adams
;                   Owen White
;                   Hamilton O. Smith
;                   J. Craig Venter
;        TITLE OF INVENTION: The Nucleotide sequence of
;                            the Haemophilus influenzae Rd Genome, Fragments
;                            Thereof, and Uses Thereof
;        NUMBER OF SEQUENCES: 1
;        CORRESPONDENCE ADDRESS:
;             ADDRESSEE: Human Genome Sciences, Inc.
;             STREET: 9410 Key West Avenue
;             CITY: Rockville,
;             STATE: MD
;             COUNTRY: USA
;             ZIP: 20850
;        COMPUTER READABLE FORM:
;             MEDIUM TYPE: 3 1/2 inch diskette
;             COMPUTER: Dell Pentium
;             OPERATING SYSTEM: MS DOS v6.22
;             SOFTWARE: ASCII Text
;        CURRENT APPLICATION DATA:
;             APPLICATION NUMBER: US/09/643,990A
;             FILING DATE: 23-Aug-2000
;             CLASSIFICATION: <Unknown>
;        PRIOR APPLICATION DATA:
;             APPLICATION NUMBER: 08/487,429
;             FILING DATE: 1995-06-07
;             APPLICATION NUMBER: 08/426,787
;             FILING DATE: 1995-04-21
;        ATTORNEY/AGENT INFORMATION:
;             NAME: Kenley K. Hoover
;             REGISTRATION NUMBER: 40,302
;             REFERENCE/DOCKET NUMBER: PB186P1C1
;        TELECOMMUNICATION INFORMATION:
;             TELEPHONE: 301-610-5790
;             TELEFAX: 310-309-8439
;   INFORMATION FOR SEQ ID NO: 1:
;        SEQUENCE CHARACTERISTICS:
;             LENGTH: 1830121 base pairs
;             TYPE: nucleic acid
;             STRANDEDNESS: double
;             TOPOLOGY: linear
;        SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-643-990A-1
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  Query Match             3.0%;  Score 99.8;  DB 4;  Length 1830121;
  Best Local Similarity   52.7%;  Pred. No. 1.9e-16;
  Matches  240;  Conservative    0;  Mismatches  212;  Indels    3;  Gaps    1;

Qy       1984 TCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTGGCAATG 2043
Db    1670240 TCTTCAGTATTTTAATGTTATTAACCGCTTGTGCGATGGCATTTGCGCACGGTTCTAATG 1670299

Qy       2044 ACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTA---TATTTGGTTTATGACACAGGAG 2100
Db    1670300 ACGTAGCCAATGCAATTGGCCCTCTTTCTGCAGTGGTCTCTATTGTAAATGAAGGCGGTA 1670359

Qy       2101 ATGTTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATCT 2160
Db    1670360 AAATTGTTTCAGGTGGAGCCTTAACTTGGTGGATTTACCCTTAGGTGCATTAGGTATTG 1670419

Qy       2161 GTGTTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACAC 2220
Db    1670420 CTGTGGGCTTAATTACTATGGGACAAAAAGTGATGGCGACTGTTGGATCTGGGATCACCG 1670479

Qy       2221 CGATCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATTG 2280
Db    1670480 ATTTAACGCCAAGCCGTGGCTTTGCCGCTCAATTTGCTACTGCGATGACCGTTGTCGTGG 1670539

Qy       2281 CATCAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGTCTG 2340
Db    1670540 CATCAGGCACAGGCTTACCCATCTCAACAACACAAACACTTGTTGGAGCTATTTTAGGTA 1670599

Qy       2341 TTGGCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTATGG 2400
Db    1670600 TCGGCTTTGCACGTGGTATCGCAGCACTGAATTTAACGGTTATCCGAAACATCATTAGTT 1670659

Qy       2401 CCTGGTTTGTCACAGTCCCCATTTCTGGAGTTATC 2435
Db    1670660 CTTGGATTGTCACATTACCAGCAGGTGCATTTTTC 1670694
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RESULT 12
US-09-252-991A-6471
; Sequence 6471, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 6471
; LENGTH: 1380
; TYPE: DNA
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-6471

```
  Query Match             2.6%;  Score 87;  DB 4;  Length 1380;
  Best Local Similarity   49.2%;  Pred. No. 7.5e-15;
  Matches  262;  Conservative    0;  Mismatches  265;  Indels    6;  Gaps    1;

Qy        514 TGCTCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCTCCGTGGGAGCCAATGATG 573
Db         78 TCCTTGTACTCGCCTGCCTGTTCGGTTTCTTCATGGCCTGGGGCGTCGGCGCCAACGACG 137

Qy        574 TAGCAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAAGCAAGCCTGCA 633
Db        138 TCGCCAACGCCATGGGCACCTCGGTGGGCTCACGGGCGCTGACCATCCGCCAGGCGATCC 197

Qy        634 TCCTAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAA 693
Db        198 TGGTGGCGATGGTCTTCGAGTTCTGCGGTGCCTACTTCGCCGGCGGCGAGGTCACCGAAA 257

Qy        694 CCATCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGGGCTACTGATGG 753
Db        258 CGATCAGGAGCGGCAT------CGTCGATCCGCAGTACATGACCCCCGACCTGCTGGTCT 311

Qy        754 CCGGCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGA 813
Db        312 TCGGCATGCTCTCGGCCTTGCTGGCCGCCGGGGTCTGGCTGTTGCTGGCGACGATCAAGG 371

Qy        814 AGCTCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGG 873
Db        372 GCTGGCCGGTGTCGACCACCCACTCCATCGTCGGCGCGATCATCGGCTTCGCCTCGGTCG 431

Qy        874 CAAAGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGATGTCTTGGTTCG 933
Db        432 GTGTCTCGGTGCACGCGGTGCACTGGGGCGCGATCGGCCCGATCGTCGCCAGTTGGGTGG 491

Qy        934 TGTCCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGTTCGTGCATTCA 993
Db        492 TGACGCCGGTGCTGTCCGGGCTGCTCGCCTTCGCCCTGTTCAGGAGCGTGCAGTGGTTGG 551

Qy        994 TCCTCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTTTC 1046
Db        552 TACTGCATGCCGAGGACCCGTTCCGCAGCGCGCGCCGCTACGTGCCGCTGTAC 604
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RESULT 13
US-09-252-991A-6303
; Sequence 6303, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 6303
; LENGTH: 1407
; TYPE: DNA
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-6303

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  Query Match             2.6%;  Score 87;  DB 4;  Length 1407;
  Best Local Similarity   49.2%;  Pred. No. 7.6e-15;
  Matches  262;  Conservative    0;  Mismatches  265;  Indels    6;  Gaps    1;

Qy        514 TGCTCATCCTGGGCTTCATTATTGCATTTGTCTTGGCATTCTCCGTGGGAGCCAATGATG 573
Db        170 TCCTTGTACTCGCCTGCCTGTTCGGTTTCTTCATGGCCTGGGGCGTCGGCGCCAACGACG 229

Qy        574 TAGCAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAAGCAAGCCTGCA 633
Db        230 TCGCCAACGCCATGGGCACCTCGGTGGGCTCACGGGCGCTGACCATCCGCCAGGCGATCC 289

Qy        634 TCCTAGCTAGCATCTTTGAAACAGTGGGCTCTGTCTTACTGGGGGCCAAAGTGAGCGAAA 693
Db        290 TGGTGGCGATGGTCTTCGAGTTCTGCGGTGCCTACTTCGCCGGCGGCGAGGTCACCGAAA 349

Qy        694 CCATCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGGGCTACTGATGG 753
Db        350 CGATCAGGAGCGGCAT------CGTCGATCCGCAGTACATGACCCCCGACCTGCTGGTCT 403

Qy        754 CCGGCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGCTTCGTTTTTGA 813
Db        404 TCGGCATGCTCTCGGCCTTGCTGGCCGCCGGGGTCTGGCTGTTGCTGGCGACGATCAAGG 463
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Qy        814 AGCTCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTGG 873
              ||  | ||    ||||| |  || ||.|| || |  || || ||| ||   || |
Db        464 GCTGGCCGGTGTCGACCACCCACTCCATCGTCGGCGCGATCATCGGCTTCGCCTCGGTCG 523

Qy        874 CAAAGGGGCAGGAGGGTGTCAAGTGGTCTGAACTGATAAAAATTGTGATGTCTTGGTTCG 933
              |  | | |  ||  | |||   |   |        || ||       |||| | |
Db        524 GTGTCTCGGTGCACGCGGTGCACTGGGGCGCGATCGGCCCGATCGTCGCCAGTTGGGTGG 583

Qy        934 TGTCCCCACTGCTTTCTGGAATTATGTCTGGAATTTTATTCTTCCTGGTTCGTGCATTCA 993
              || | ||  |||| || ||  | |  |       | |||       || |      ||
Db        584 TGACGCCGGTGCTGTCCGGGCTGCTCGCCTTCGCCCTGTTCAGGAGCGTGCAGTGGTTGG 643

Qy        994 TCCTCCATAAGGCAGATCCAGTTCCTAATGGTTTGCGAGCTTTGCCAGTTTTC 1046
              | || |||   |  || ||  | |  |  |     ||    ||||  | | |
Db        644 TACTGCATGCCGAGGACCCGTTCCGCAGCGCGCGCCGCTACGTGCCGCTGTAC 696


RESULT 14
US-09-198-452A-1/c
; Sequence 1, Application US/09198452A
; Patent No. 6559294
; GENERAL INFORMATION:
;  APPLICANT: Griffais, R.
;  TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
;  TITLE OF INVENTION:  thereof and uses thereof, in particular for the diagnosis, preve
;  TITLE OF INVENTION:  and treatment of infection
;  FILE REFERENCE: 9710-003-999
;  CURRENT APPLICATION NUMBER: US/09/198,452A
;  CURRENT FILING DATE:  1998-11-24
;  NUMBER OF SEQ ID NOS: 6849
; SEQ ID NO 1
;   LENGTH: 1230025
;   TYPE: DNA
;   ORGANISM: Chlamydia pneumoniae
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)..(15000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (15001)..(30000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (30001)..(45000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (45001)..(60000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (60001)..(75000)
;   OTHER INFORMATION: n=a or c or g or t
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;   LOCATION: (120001)..(135000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
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;   OTHER INFORMATION: n=a or c or g or t
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;   LOCATION: (150001)..(165000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (165001)..(180000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (180001)..(195000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
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;   NAME/KEY: misc_feature
;   LOCATION: (210001)..(225000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
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;   LOCATION: (240001)..(255000)
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;   LOCATION: (255001)..(270000)
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;   NAME/KEY: misc_feature
;   LOCATION: (270001)..(285000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (285001)..(300000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (300001)..(315000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (315001)..(330000)
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;   NAME/KEY: misc_feature
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;   NAME/KEY: misc_feature
;   LOCATION: (390001)..(405000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (405001)..(420000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (420001)..(435000)
;   OTHER INFORMATION: n=a or c or g or t
;   NAME/KEY: misc_feature
;   LOCATION: (435001)..(450000)
;   OTHER INFORMATION: n=a or c or g or t
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Qy       1984 TCTTCCAGTTCCTGCAGATCCTTACAGCCTGCTTTGGGTCATTCGCCCATGGTGGCAATG 2043
             ||||     | ||| |||||  |   ||| ||||||  | | || || || ||    ||||
Db     773577 TCTTTGCCTACCTACAGATTATCGTAGCTTGCTTTATGGCGTTTGCTCACGGATCTAATG 773518

Qy       2044 ACGTAAGCAATGCCATTGGGCCTCTGGTTGCTTTATATTTGGTTTATGACACAGGAGATG 2103
             | ||    |||||||||  ||| | | ||   | |       |   | |  ||      ||
Db     773517 ATGTTGCTAATGCCATTGCTCCTGTAGCTGGAGTCT------TGCGTCAGGCATATCCTG 773464

Qy       2104 TTTCTTCAAAAGTGGCAACACCAATATGGCTTCTACTCTATGGTGGTGTTGGTATCTGTG 2163
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Db     773463 CTTCCTATACGTCGTATACATTAATTAGGCTCATGGCATTTGGAGGCATAGGCTTGGTCA 773404

Qy       2164 TTGGTCTGTGGGTTTGGGGAAGAAGAGTTATCCAGACCATGGGGAAGGATCTGACACCGA 2223
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Qy       2224 TCACACCCTCTAGTGGCTTCAGTATTGAACTGGCATCTGCCCTCACTGTGGTGATTGCAT 2283
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Qy       2284 CAAATATTGGCCTTCCCATCAGTACAACACATTGTAAAGTGGGCTCTGTTGTGTCTGTTG 2343
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Qy       2344 GCTGGCTCCGGTCCAAGAAGGCTGTTGACTGGCGTCTCTTTCGTAACATTTTTATGGCCT 2403
              | |     ||    |     ||  || |||       |  |       | |||  |  |||
Db     773223 GTTTAGCACGAGGGATCCGTGCCATTAACTTAAACATTATCAAAGATATTGTACTCTCCT 773164

Qy       2404 GGTTTGTCACAGTCCC 2419
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Db     773163 GGTTTATTACGCTTCC 773148
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RESULT 15
US-09-103-840A-2
; Sequence 2, Application US/09103840A
; Patent No. 6294328
; GENERAL INFORMATION:
;  APPLICANT: FLEISCHMAN, Robert D.
;  APPLICANT:  WHITE, Owen R.
;  APPLICANT:  FRASER, Claire M.
;  APPLICANT:  VENTER, John C.
;  TITLE OF INVENTION: DNA SEQUENCES FOR STRAIN ANALYSIS IN MYCOBACTERIUM
;  TITLE OF INVENTION:  TUBERCULOSIS
;  FILE REFERENCE: 24366-20007.00
;  CURRENT APPLICATION NUMBER: US/09/103,840A
;  CURRENT FILING DATE:  1998-06-24
;  NUMBER OF SEQ ID NOS: 2
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; SEQ ID NO 2
;   LENGTH: 4403765
;   TYPE: DNA
;   ORGANISM: Mycobacterium tuberculosis
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;   OTHER INFORMATION: CDC 1551
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;   OTHER INFORMATION: represent a, t, c or g
US-09-103-840A-2
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Qy        573 GTAGCAAATTCTTTTGGTACAGCTGTGGGCTCAGGTGTAGTGACCCTGAAGCAAGCCTGC 632
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Qy        693 ACCATCCGGAAGGGCTTGATTGACGTGGAGATGTACAACTCGACTCAAGGGCTACTGATG 752

Db    2549366 ACCATCCGCAGCGGCATCGTTGATCTGTCCGGGGTGTCCGTCGACCCACGCGACTTCATG 2549425

Qy        753 GCCGGCTCAGTCAGTGCTATGTTTGGTTCTGCTGTGTGGCAACTCGTGGCTTCGTTTTTG 812

Db    2549426 AACATCATGCTGTCGGCGCTATCGGCAGCCGCGCTCTGGCTGCTGTTTGCTAACCGTATG 2549485

Qy        813 AAGCTCCCTATTTCTGGAACCCATTGTATTGTTGGTGCAACTATTGGTTTCTCCCTCGTG 872

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Job time : 206 secs